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In-situ measurement of the effect of canopy tree fine roots on nitrogen availability in forest soil

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(Declaration of interest)

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Abstract

Plant fine roots affect the amount of soil inorganic nitrogen (N) via their N uptake and their effects on organic matter decomposition and N mineralization. However, the effects of the fine roots of canopy trees on N availability in forest soil remain largely unknown owing to the limitations of *in-situ* measurements. Therefore, to reveal the effects of the fine roots of canopy trees on forest soil N availability, we developed a novel *in-situ* method that combines a modified *in-situ* resin-core method, using live fine roots, with a tree N demand estimation, using the N-balance method. Furthermore, we measured soil extracellular enzyme activities in the rhizosphere and bulk soils to determine whether fine roots stimulate soil enzyme activities. The results showed that soil enzyme activities were stimulated by the fine roots of canopy trees, especially in the rhizosphere, and this indicates that fine roots stimulate organic matter decomposition and N mineralization. However, fine roots had no significant effect on either the pool size or leaching of inorganic N. The estimated amount of potential N uptake by fine roots was greater than the amount of inorganic N released via fine root-induced N mineralization. Overall, our results indicate that almost all of the fine root-induced increment of mineralized inorganic N is taken up by the fine roots; thus, the fine roots of canopy trees did not affect N availability in forest soil. This study was limited by the fact that the fine roots in the resin cores were less dense than natural roots, and the fact that the actual root N uptake can differ from the estimated N uptake; these limitations should be resolved in future studies. Nevertheless, the proposed

19 method can be used to investigate the effects of fine roots on the N cycle in forest soil under field

20 conditions considering N uptake and priming.

21 **Keywords:** Nitrogen cycle, Tree nitrogen uptake, Rhizosphere priming, Extracellular enzyme

22 activities

23

Introduction

Nitrogen (N) availability in forest soils often limits the primary production of temperate forests (Vitousek and Howarth, 1991; LeBauer and Treseder, 2008). Most soil N is bound in complex macromolecular soil organic matter (SOM), a form of N that plants cannot directly utilize in forest ecosystems (Bormann et al., 1977). Complex SOM is degraded into labile SOM through the activities of various enzymes; labile SOM is further decomposed by enzymes and finally mineralized into inorganic N (Geisseler et al., 2010). Because net N mineralization is significantly and positively correlated with the annual net primary production of forests (Reich et al., 1997), understanding N mineralization processes is key for understanding forest productivity.

Plant roots not only affect soil N pools through their uptake of soil N but also by altering the activities of enzymes involved in SOM decomposition via rhizodeposition (Brzostek et al., 2013; Yin et al., 2014; Meier et al., 2017), and this consequently affects the N mineralization rate in the rhizosphere (i.e., the soil surrounding plant roots). This rhizodeposition-induced alteration of the decomposition and mineralization processes is called the rhizosphere priming effect (RPE) and has been measured mainly through the changes in soil respiration (Dijkstra et al., 2013). The RPE ranges from -50% to 380% and is influenced by factors such as plant type, plant growth stage, and environmental conditions, including soil temperature and nutrient conditions (Cheng et al., 2003,

2014; Cheng and Kuzyakov, 2005; Dijkstra and Cheng, 2007; Bengtson et al., 2012). Tree species tend to have a higher level of RPE than grassland or crop species (Yin et al., 2014; Huo et al., 2017). Furthermore, most trees in temperate forests have symbiotic relationships with mycorrhizal fungi (Marschner and Dell, 1994; Deckmyn et al., 2014). Besides helping their host plants to take up nutrients, including N, mycorrhizal fungi deliver photosynthetically produced organic carbon (C) from the host to other parts of the fungus via their hyphae, which promotes the degradation of organic N that is far from the rhizosphere (Marschner and Dell, 1994; van Hees et al., 2006; Toljander et al., 2007; Talbot and Treseder, 2010). Therefore, mycorrhizal symbiosis would extend the priming effects of plant roots into not only the rhizosphere but also the bulk soil. However, most of these findings are based on studies of the priming effect on C dynamics, rather than N dynamics. Although studies using herbaceous species and seedlings and saplings of woody species have shown that the priming effect also affects soil N availability (Dijkstra et al., 2009; Bengtson et al., 2012; Cheng et al., 2014), there is relatively limited information on the priming effects of the fine roots of canopy trees on forest soil N dynamics owing to methodological limitations.

Specifically, nitrogen mineralization processes, including ammonification and nitrification, in forest soils have been investigated mainly using laboratory incubation methods and *in-situ* incubation methods (Kandeler, 1996). For *in-situ* incubations, the buried bag and resin core methods have been widely used (e.g. Hart and Firestone 1989; Zou et al. 1992; Kolberg et al. 1997; Shibata et

al. 2011; Urakawa et al. 2014). Studies using these methods have revealed that N transformation is affected by environmental factors, such as temperature, moisture content, and C and N concentrations (Kladivko and Keeney, 1987; Tietema et al., 1992; Yan et al., 2008; Shibata et al., 2011). However, as these traditional methods exclude the effects of live plant roots, they may overestimate or underestimate N transformation processes (Frank and Groffman, 2009). Furthermore, studies on the priming effects of the fine roots of canopy trees on the N cycle have predominantly compared the mineralization potential of field-sampled rhizosphere soil to that of bulk soil using short-term laboratory incubations without live roots (Brzostek et al., 2013; Yin et al., 2014; Lin et al., 2018). Comparisons made using this technique reveal the potential RPE on N mineralization; however, the method still ignores the organic substrate supply and nutrient uptake by fine roots and mycorrhiza during incubation. Nitrogen diffusion according to concentration gradients and N loss through leaching are also not considered in the laboratory incubation method. Therefore, *in-situ* measurements with live roots are necessary to understand the potential effects of the fine roots of canopy trees on the N cycle in forest soils under natural conditions.

Here, we aim to reveal how canopy tree roots affect forest soil N dynamics via uptake and priming. For this purpose, we developed a new method that combines *in-situ* incubation and root N uptake estimation. For *in-situ* incubation, we modified the resin core method by inserting an intact root system of a canopy tree into the side of an unplasticized polyvinyl chloride (uPVC) pipe (Fig. 1).

Thereafter, the changes in the extractable N content in the incubated soil and N leaching from the bottom of the pipe were measured; net inorganic N increment was defined as the total increase in inorganic N pools and leaching. To estimate the potential N uptake by fine roots in the plots and pipes, the above- and belowground net primary production were calculated based on litterfall, fine root production, and the biomass growth estimated using allometric relationships. The amount of change in the inorganic N pools change that was induced by the fine roots of canopy trees was evaluated by comparing the net inorganic N increments and potential N uptakes between incubations conducted with and without live roots. As the RPE and hyphosphere priming effect vary with nutrient availability (Phillips and Fahey, 2008), we investigated the differences in the magnitude of the effects of fine roots on soil N availability between soil with and without an organic horizon. The activities of four enzymes were measured at the end of the incubation to assess the positive effects that rhizosphere and hyphosphere priming had on N mineralization through their stimulation of SOM decomposition. Studies have reported that tree roots significantly accelerate nutrient cycling, and that this acceleration is quantitatively important (Finzi et al., 2015; Huo et al., 2017). At the same time, clear-cutting of trees increases ecosystem N losses (Vitousek et al., 1979), indicating that trees increase N retention in the forest ecosystem through their uptake and storage of N in their biomass. Therefore, we hypothesize that (1) the fine roots of canopy trees will stimulate enzyme activities, resulting in the stimulation of N mineralization via enhanced SOM decomposition. However, (2) the amount of N taken up by fine

roots will exceed the amount of N mineralized as a result of priming effects. Therefore, (3) fine roots will reduce the size of the inorganic N pools within the pipes and the extent of N leaching from the bottom of the pipes, indicating that canopy trees increase N retention in the forest ecosystem.

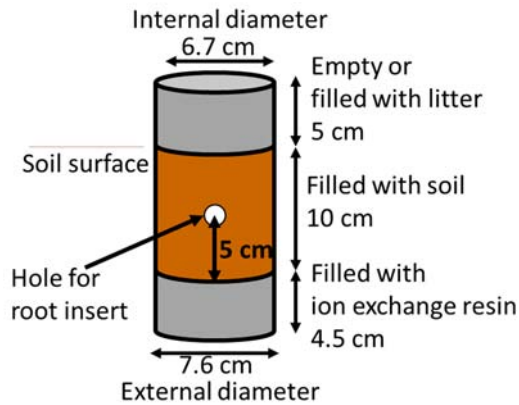


Fig. 1 Schematic of the incubation pipe used in this study.

Experimental

Study site

The study site was a deciduous broad-leaved temperate forest in the Shibeche Branch of the Hokkaido Forest Research Station at the Field Science Education and Research Center of Kyoto University (43°24.2'N, 144°38.5'E). The soil in this site has been characterized as an Andosol (IUSS Working Group WRB, 2015). According to data from a meteorological station located approximately 9 km south of the study site, the mean annual precipitation and air temperature are 1189 mm and 6.3 °C

(1986–2015), respectively. Soil freezing occurs from December to May, and the soil is covered by snow from November to April (Hosokawa et al., 2017).

We established four plots (15 m × 15 m) in the pure forest of *Quercus crispula* Blume. Each plot was separated from other plots by at least 30 m. The only canopy tree species in the study plots was *Q. crispula*, and the understory of the forest stand was densely covered with dwarf bamboo (*Sasa nipponica* Makino et Shibata). *Quercus crispula* has symbiotic relationships with ectomycorrhizal and arbuscular mycorrhizal fungi (Obase et al., 2007) and is a common tree species in the cool temperate forests of Japan (Hiura, 2001; Ohsawa et al., 2011). The number of *Q. crispula* in the study plots was 7–10, and its root density at the depth of 0–10 cm was $132.9 \pm 47.4 \text{ g m}^{-2}$ (Nakayama and Tateno, 2018). Most of the fine roots were distributed in the soil at depths of 0–50 cm (Gale and Grigal, 1987), and the vertical fine root productivity decreased with increasing soil depth, as did the fine root density (Fukuzawa et al., 2007). Therefore, almost all N for *Q. crispula* was taken up by the fine roots located within the top 50 cm of the soil. As 53.7% of the fine roots in the top 50 cm of the soil were concentrated at a depth of 0–10 cm in the adjacent plot (Tateno et al., 2020), the fine root density at the depth of 0–50 cm in the study plot was estimated as $247.4 \pm 88.3 \text{ g m}^{-2}$. Details of study site and plots can be found in previous reports (Hosokawa et al., 2017; Nakayama and Tateno, 2018; Nakayama et al., 2019; Watanabe et al., 2019; Tateno et al., 2020).

In-situ incubation

Before incubation, the volumes of the mineral soil and the organic horizon used in the incubation were determined by sampling two surface soil samples (0–10 cm-depth) and one organic horizon from each plot on May 28, 2016; mineral soil was sampled using a soil core (surface area; 20 cm²) and the organic horizon was manually collected from a 40 cm × 40 cm area. The soil samples were passed through a 4-mm mesh sieve to remove fine roots, gravel, and organic material. The organic horizon was divided into three groups (able to pass through a 9 mm mesh sieve, larger than 9 mm and partially decomposed, and larger than 9 mm and relatively fresh). After sieving and separating, the total weights of the mineral soil and organic horizon samples were measured. Thereafter, each sample was divided, and a subsample from each sample was oven-dried (at 70 °C for more than 72 h) to measure its moisture content. To calculate the water content of each wet soil and organic horizon sample, we multiplied the wet weight of the entire sample by the moisture content of its oven-dried subsample. Then, the bulk density and total dry weight of the organic horizon in the plots was calculated by subtracting the weight of the water contents from that of the total wet soil and organic horizon.

Soil samples (from the 0–10-cm depth) for the incubation were collected on June 7, 2016 from four randomly selected points in each plot and pooled to form one composite sample per plot. The soil samples were passed through a 4 mm mesh sieve and stored at 4 °C until further treatment. A

part of the newly collected soil was oven-dried at 70 °C for 72 h to measure the moisture content. Based on the above-mentioned soil bulk density and moisture content of the newly collected soil, we determined the soil (wet) weight to be used in the incubation. The bulk density of soil in the incubation pipes was the same as that of the surrounding soil. We also determined the weights of the three groups of sieved organic horizon material for incubation and used amounts that were consistent with the dry weight of the plots on a per area basis. Some of the soil samples were used to measure the initial soil extractable N concentration, as described below. The pH of the soil used in the incubation was 5.0 ± 0.1 . The total C and N contents of the mineral soil and organic horizon used in the incubation are described in Table 1.

The prepared soils were incubated *in-situ* using an altered resin-core method with uPVC pipes (VP65; Sekisui chemical CO., LTD, Osaka, Japan; inner diameter: 6.7 cm, height: 15 cm; Fig. 1). We pierced a hole on the side of half of the pipes for the roots (diameter: 8 mm; the center of the hole was 5 cm from the bottom of the pipe). Short uPVC pipes (height: 4.5 cm) were filled with 45 g wet weight of a mixed bed ion-exchange resin (MB-1; ORGANO CORPORATION, Tokyo, Japan) and covered with nylon mesh to retain the resin. The short pipes were placed at the bottom of all incubation pipes to capture the inorganic N leached during the incubation period (Fig. 1). Soil was prevented from escaping from the bottom of the incubation pipes by the nylon mesh. Sixty-four pipes (32 pierced and 32 non-pierced) were used. On June 9, 2016, 32 live, intact fine-root systems of *Q.*

crispula (eight root systems from each plot) were excavated at a soil depth of 0–10 cm by hand. Thereafter, 12.5 cm-deep holes were dug, taking care not to harm the excavated roots, and the pierced pipes were inserted into the holes. The live intact root system (approximately 15 cm-long) was carefully inserted into each pierced pipe through the hole in the pipes. The holes for root insertion were then sealed with silicone sealant. For the 32 non-pierced pipes, we dug 12.5 cm-deep holes and inserted the pipes into the holes without the live root systems. Soil was added and pushed gently from the top of the pipes to fill 0–10 cm of the pipes and ensure that the bulk density of soil for incubation was the same as that of the surrounding soils. Subsequently, we placed the sampled organic horizon on top of the soils in half of the pipes with root systems and half of the pipes without root systems, ensuring the same density and composition as the surrounding organic layer. The tops of the pipes were covered with a rough polyester drainage net to prevent litter input during incubation, while still allowing water flow from the upper surfaces of the pipes. Thus, four treatments were analyzed in this study: with root system and organic horizon (RL), with root system but without organic horizon (R), without root system but with organic horizon (CL), without root system or organic horizon (C). Four replicates for each plot and treatment were incubated, with a total of 16 pipes for each plot and treatment.

All incubation pipes were collected on June 5, 2017 (after an incubation period of 361 d).

We defined the soil adhering to the roots after gentle shaking as rhizosphere soil (Phillips and Fahey,

2006, 2008). Samples of bulk and rhizosphere soils were mixed for each treatment and plot. The roots incubated in two RL samples in plot 2 and one RL sample and two R samples in plot 3 were assumed to have died during the incubation based on their color and morphology (black and soft). Therefore, we did not mix them into the composite samples. After mixing, some of the bulk soil samples were oven-dried (70 °C, 72 h) to measure the moisture content. Because the amount of rhizosphere samples was limited, we assumed that the moisture content of rhizosphere samples was the same as that of bulk soils. The roots from each pipe were washed, oven-dried (70 °C, 72 h), and weighed.

Measurements of soil N concentrations

Pre-incubation wet soils and wet bulk soil samples were used to determine NO_3^- -N (nitrate), NH_4^+ -N (ammonium), and dissolved organic N (DON) concentrations after extraction with 0.5 M K_2SO_4 (extracted from 5 g of wet sample into 50 mL of K_2SO_4). Ion-exchange resin (10 g, wet weight) placed at the bottom of the incubation pipes was extracted twice with 50 mL of 1 M KCl. Ammonium and nitrate concentrations were measured by colorimetry using a microplate reader (Synergy HXT; BioTek, Winooski, VT, USA) with the indophenol blue method and Griess assay (Miranda et al., 2001). To measure the total extractable N (TEN), 1 mL extracts were autoclaved (121 °C, 20 min) with 1 mL of ion-exchanged water and 200 μL of alkalinity potassium persulfate solution. Thereafter, TEN was

measured using the same method used for nitrate measurements. The DON concentration of each sample was calculated by subtracting the summed of inorganic N (i.e., nitrate + ammonium) concentration from the TEN concentration. To measure the microbial biomass N (MBN) concentration, 10 g of wet soil was fumigated with chloroform. After 24 h of fumigation, the TEN content of fumigated soil was measured in the same manner described above. The MBN concentration was multiplied by 2.2 after subtracting the TEN concentration of the non-fumigated extracts from that of the fumigated extracts (Inubushi et al., 1984).

Measurement of soil extracellular enzyme activities

The enzyme assays were conducted within 24 h of sampling. The activities of four extracellular enzymes (Table S1) were measured using wet bulk and rhizosphere soil samples following the method of Saiya-Cork et al. (2002). Briefly, wet soil samples (2.0 g for bulk soil, 1.0 g for rhizosphere soil) were suspended in 200 mL (bulk soil) or 100 mL (rhizosphere soil) of acetate buffer (50 mM sodium acetate; pH 5.0). The soil suspensions were homogenized using a magnetic stirrer for 1 min. Subsequently, 200 μ L aliquots of the suspensions were dispensed into 96-well microplates (eight replicate wells per sample per assay). Fifty microliters of a substrate (200 μ M) was added into the sample wells. In the blank well, 50 μ L of acetate buffer and 200 μ L of sample suspension were

dispensed. Fifty microliters of 10 μ M 4-methylumbelliferone (MUB) was added into the quenching standard well with 200 μ L of sample suspension. The reference standard well had 200 μ L of acetate buffer and 50 μ L of 10 μ M 4-MUB. The negative control well had 200 μ L of acetate buffer and 50 μ L of the substrate. The microplates were incubated at 20 °C in the dark for up to 2 h, and then the fluorescence was measured using a microplate reader (Synergy HXT; BioTek) with excitation and emission wavelengths of 360 and 460 nm, respectively. The target enzymes, incubation times, and substrates are listed in Table S1.

Quantification of net primary production and tree N demand

To estimate the net primary production of *Q. crispula*, litterfall and fine root growth were measured, and woody biomass (trunk, branch, and coarse root) production was estimated using the diameter at breast height (DBH) and the allometric relationships. To collect litterfall, eight litter traps (two for each plot; opening area: 0.25 m²) were placed in the study plots. Litterfalls were collected once a month from September until all leaves of *Q. crispula* had fallen (September 26, October 18, and November 10, 2016). In the laboratory, the collected litter was oven-dried (at 60 °C for more than 72 h). We separated the dried litter into five categories (leaves of *Q. crispula*, leaves of other plants, small branches, large branches, and others) by hand sorting, and then weighed the litter. Others in the litter

trap (hereafter; other litter) included the acorns, flowers, and bud scales of *Q. crispula*. Fine root growth was measured using the ingrowth core method. On July 5, 2016, eight cylindrical ingrowth cores (mesh size: 2 mm, inner diameter: 3 cm, height: 10 cm) were buried in each plot (total 32 cores). The cores were collected on August 15, 2017 (incubation period: 406 days). The cores contained roots of *Q. crispula* and *S. nipponica*, which were manually separated according to their morphological traits including color and their associated mycorrhiza. Both types of root were washed with water, oven-dried (70 °C, 72 h), and weighed. The annual fine root growth of *Q. crispula* (0–50 cm depth) was estimated from the fine root dry weight of *Q. crispula* in the ingrowth cores (0–10 cm) and vertical fine root distribution (0–50 cm) in the adjacent plot, assuming that fine root growth is proportional to fine root biomass. In other words, the production of fine roots of *Q. crispula*, as observed in ingrowth cores, was divided by the incubation period and the proportion of fine root of *Q. crispula* in the 0–10-cm soil layer of the adjacent plot (53.7%). To estimate woody biomass production, we measured the DBH of each tree in the plot on July 13, 2016 and June 8, 2020. Two individuals died during 2016–2020; therefore, the growth of these trees was assumed as zero. The growth of some individuals that exhibited negative growth of DBH due to bark peeling was also assumed as zero. We used the following allometric relationships for *Q. crispula* (Takagi et al., 2010):

$$Y_t = e^{(2.365 \times \log_e DBH - 2.596)}$$

$$Y_b = e^{(2.713 \times \log_e DBH - 4.456)}$$

$$Y_c = e^{(2.224 \times \log_e DBH - 2.918)}$$

where, Y_t , Y_b , and Y_c represent the biomass of the trunk, branch, and coarse root (kg), respectively. These values were divided by four, which was the interval of DBH measurement (years), to estimate the annual production. The annual net primary production ($\text{g m}^{-2} \text{ year}^{-1}$) was estimated as the sum of leaf litter, other litter, the estimated annual fine root growth (0–50 cm), and the estimated production of the trunks, branches, and coarse roots of *Q. crispula*.

To estimate the annual N demand of *Q. crispula* (N_{demand}), we calculated its total N content as the sum of the N contents of its leaf litter, other litter, fine roots, and woody parts (Aber et al., 1985; Finzi et al., 2007; Tateno and Takeda, 2010; Lü et al., 2014). Before measuring the total N concentration, leaf litter collected using the litter traps was mixed for each trap and species ($n = 8$), and the fine roots collected from the ingrowth cores were mixed for each plot and species ($n = 4$). Because the amount of other litter was limited, we composited all of other litter samples into one before next step. Leaf litter, other litter, and fine roots were ground before measuring their total N concentrations. The total N concentrations of the leaf litter, other litter, and roots were measured using an elemental analyzer (Sumigraph NC-900; Sumika Chemical Analysis Service, Ltd.). The total N concentrations of the branches, trunks, and coarse roots of *Q. crispula* were estimated as 0.58%, 0.39%, and 0.45%, respectively (Takagi et al., 2010). The N_{demand} ($\text{g-N m}^{-2} \text{ year}^{-1}$) of *Q. crispula* was calculated by multiplying the annual net primary production by the N concentration in each part of *Q.*

crispula (Table 2). The potential N uptake per gram of dry *Q. crispula* fine roots in the plots under natural conditions (N_{uptake} ; $\text{mg-N g}^{-1} \text{ dry root year}^{-1}$) was estimated by dividing N_{demand} by the density of *Q. crispula* fine roots in the plots (0–50 cm depth) (Table 2).

Data calculation and statistical analysis

The annual inorganic N increment (N_{inc} ; $\text{g-N m}^{-2} \text{ year}^{-1}$) and annual nitrate increment (Nit_{inc} ; $\text{g-N m}^{-2} \text{ year}^{-1}$) were calculated based on the ammonium and nitrate pools and leaching, as listed in Table 2. Because N inputs from rain and leachates of the organic horizon entering from the tops of the pipes were not excluded to make the conditions inside the pipes as natural as possible, these N inputs were included in N_{inc} and Nit_{inc} . The root-induced annual inorganic N increment under the conditions in the incubation pipes (N_{induced} ; $\text{g-N m}^{-2} \text{ year}^{-1}$) was also estimated (Table 2). According to some reports, nitrogen uptake by plants varies with soil N availability (e.g., Liao et al., 2018), whereas other studies have reported the N availability is not significantly related to N uptake (e.g., Tateno and Takeda, 2010; Maeda et al., 2018). Therefore, the “actual” N uptake by fine roots could vary with the environmental conditions. However, to estimate the amount of potential N uptake in incubation pipes, we assumed that the potential N uptake per unit fine root dry mass (g) in the incubation pipes was the same as that under natural conditions. The potential amount of N taken up by the fine roots of *Q. crispula* under

the conditions in the incubation pipe (N_{uptake} ; $\text{g-N m}^{-2} \text{ year}^{-1}$) was calculated by multiplying N_{at} N_{uptake} by the fine root density in each incubation pipe (Table 2).

The difference in the root densities within the pipes at the end of the incubation between the two root treatments (R vs. RL) was tested using a t-test. The changes in the N concentrations during the incubation were tested using a one-way analysis of variance (ANOVA) with a random effect (random effect: sampling plot). To evaluate the differences in the soil properties (moisture content and extractable N concentrations) and the amounts of leached inorganic N among the treatments (C, CL, R, and RL), two-way ANOVAs with random effects (random effect: sampling plot) were used. To analyze the differences in the soil extracellular enzyme activities between the bulk and rhizosphere soils and among the treatments, we used a two-way ANOVA with a random effect (random effect: sampling plot). The RPEs on the activities of soil extracellular enzymes were calculated by dividing the enzyme activities in rhizosphere by those in the bulk soil. Furthermore, to compare the magnitude of the RPE on enzyme activity, a one-way ANOVA with a random effect (random effect: sampling plot) was performed for each type of enzyme examined. All statistical analyses were conducted using R software (ver. 3.5.0, R Core Team, 2018).

Results

Root density and soil moisture content at the end of incubation

The root densities of R and RL in the pipes at the end of incubation were $66.63 \pm 17.44 \text{ mg pipe}^{-1}$ and $63.06 \pm 11.15 \text{ mg pipe}^{-1}$, which were equivalent to $18.90 \pm 4.95 \text{ g m}^{-2}$ and $17.89 \pm 3.88 \text{ g m}^{-2}$ (0–10 cm-depth) for R and RL, respectively. There was no significant difference between the root densities of R and RL (t-test, $p = 0.745$). The moisture contents of the soil were $52.2 \pm 1.2\%$, $52.9 \pm 1.7\%$, $51.0 \pm 0.9\%$, and $51.7 \pm 0.8\%$ for C, CL, R, and RL, respectively. Although the variation in moisture content was only few percent, the effects of roots and organic horizon on the moisture content were statistically significant (two-way ANOVA with a random effect; $F = 18.9$, $p < 0.01$ for roots and $F = 6.1$, $p < 0.05$ for the organic horizon). The interaction effect of roots and the organic horizon on moisture content was not significant (two-way ANOVA with a random effect; $F = 0.0$, $p = 0.985$).

Changes in N concentration and differences in N concentration and leaching during incubation among the treatments

Regardless of the treatment, the soil nitrate concentration after incubation was significantly higher than that before incubation (one-way ANOVA with a random effect; $F = 12.7$, $p < 0.05$; Fig. 2a). The roots and organic horizon had no significant effects on the concentration of nitrate after incubation

(Fig. 2a). In contrast, the soil ammonium concentration after incubation was significantly lower than that before incubation (one-way ANOVA with a random effect; $F = 75.1$, $p < 0.001$; Fig. 2b). A significant effect of the roots and organic horizon interaction on the ammonium concentration after incubation was detected using a two-way ANOVA with a random effect. We then divided the samples based on presence or absence of roots and organic horizons. Among samples without root systems (C and CL), the concentration of ammonium was significantly higher under C than under CL (one-way ANOVA with a random effect; $F = 18.0$, $p < 0.05$), whereas the organic horizon did not have a significant effect on the ammonium concentration in samples with root systems (one-way ANOVA with a random effect; R vs. RL, $F = 1.99$, $p = 0.254$). After dividing the samples into those with and without organic horizons, the roots did not have a significant effect on the ammonium concentration (one-way ANOVA with a random effect; $F = 2.23$, $p < 0.233$ for CL vs. RL and $F = 3.43$, $p = 0.161$ for C vs. R). The DON and MBN concentrations did not change significantly during incubation (one-way ANOVA with a random effect; $F = 1.30$, $p = 0.273$ and $F = 0.74$, $p = 0.404$, respectively). There was no significant difference in the DON and MBN concentrations after incubation among the treatments (Fig. 2c, d).

The amount of inorganic N leached from the bottom of the incubation pipes during the incubation ranged from 3.86 to 7.92 g-N m⁻²; 96.1% ± 1.7% of the leached N was nitrate (Fig. 3). The amount of leached nitrate was significantly higher in samples with organic horizons (Fig. 3a). The

treatment type did not have a significant effect on the amount of leached ammonium (Fig. 3b).

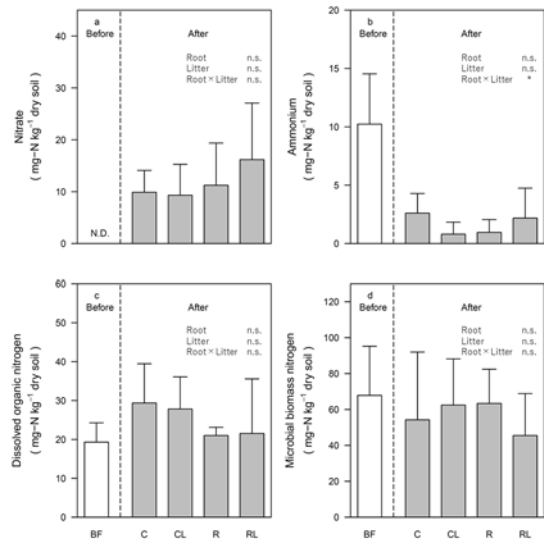


Fig. 2 Concentrations of (a) nitrate, (b) ammonium, (c) dissolved organic nitrogen, and (d) microbial biomass nitrogen before (white bars) and after (gray bars) incubation. The bars and error bars represent the mean and standard deviation, respectively (n = 4). N.D. indicates that the concentration was under the detection limit. C, CL, R, and RL represent each treatment, with R indicating treatments with root systems, C indicating those without root systems, and L indicating treatments with organic horizons (i.e., litter). BF means before incubation and represents the initial soil. Root, Litter, and Root × Litter in the upper right of each figure represent the results of the two-way ANOVA with a random effect. The symbols are as follows: n.s.: $p \geq 0.05$, *: $p < 0.05$

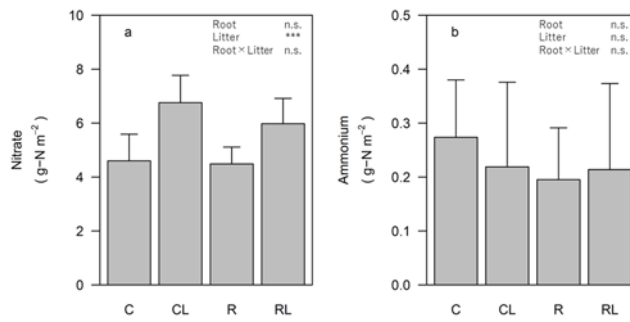


Fig. 3 Amounts of (a) nitrate and (b) ammonium leached from the bottom of the incubation pipes during the incubation. The bars and error bars represent the mean and standard deviation, respectively ($n = 4$). C, CL, R, and RL represent each treatment. Root, Litter, and Root \times Litter in the upper right of each figure represent the results of the two-way ANOVA with a random effect. The symbols are as follows: n.s.: $p \geq 0.05$, ***: $p < 0.001$

Above- and belowground primary production of *Q. crispula*

The net primary production and N concentration in each part of *Q. crispula* are shown in Table 3. Leaf production was the highest, followed by trunk, branch, and coarse-root production. The nitrogen concentration was also the highest in the leaf litter ($1.82\% \pm 0.08\%$, Table 3). Approximately three-quarters of the N_{demand} of the canopy *Q. crispula* trees was from leaf litter production (Table 3).

Potential nitrogen uptake by *Q. crispula*, N budget, and root effect on the N budget

Based on the N_{demand} and fine root density of the study plots, the estimated N_{uptake} was $28.58 \pm 5.86 \text{ mg-N g}^{-1} \text{ dry root year}^{-1}$. The N_{uptake} was estimated as $0.53 \pm 0.15 \text{ g-N m}^{-2} \text{ year}^{-1}$.

Regardless of treatment, positive N_{itinc} and N_{inc} occurred (Table 4). Samples with organic horizons presented higher N_{itinc} and N_{inc} values than those without organic horizons (two-way ANOVA with a random effect; $F = 119.5$, $p < 0.001$ and $F = 99.6$, $p < 0.001$, respectively), indicating N was leached from the organic horizon. The effect of roots was not significant (two-way ANOVA with a random effect; $F = 2.89$, $p = 0.123$ for N_{itinc} and $F = 3.33$, $p = 0.101$ for N_{inc} , respectively), whereas samples with root systems tended to have lower N_{itinc} and N_{inc} values (Table 4). The effect of the root and organic horizon interaction was not significant (two-way ANOVA with a random effect; $F = 1.79$, $p = 0.214$ for N_{itinc} and $F = 0.56$, $p = 0.473$ for N_{inc} , respectively).

The average N_{induced} was positive, however, the magnitude of the effect differed among the treatments (Table 4). The average N_{induced} in samples with the organic horizon was almost zero and lower than that in samples without the organic horizon (Table 4).

Enzyme activities in the rhizosphere and bulk soils

All enzyme activities investigated in this study were on average 2.7–4.4 times higher in the rhizosphere soils than in the bulk soils, irrespective of the treatment (Fig. 4). The RPEs on the activities of acid

phosphatase and NAGase were slightly higher in samples without organic horizons ($478\% \pm 286\%$ and $545\% \pm 313\%$, respectively) than in those with organic horizons ($391\% \pm 241\%$ and $338\% \pm 115\%$, respectively); however, the difference was not statistically significant (one-way ANOVA with a random effect; $F = 5.34$, $p = 0.104$ and $F = 1.54$, $p = 0.303$, respectively). The RPEs on the activities of glucosidase and xylosidase were similar between samples with ($253\% \pm 46\%$ and $286\% \pm 71\%$, respectively) and without ($277\% \pm 70\%$ and $244\% \pm 45\%$, respectively) organic horizons, and the difference of treatments was not significant (one-way ANOVA with a random effect; $F = 0.58$, $p = 0.502$ and $F = 3.34$, $p = 0.165$, respectively). Within the treatments with organic horizons, all enzyme activities in the bulk soils were slightly higher in the treatment with root systems than in that without root systems (Fig. 4). This trend was not consistent for the activities of acid phosphatase and NAGase in the treatments without organic horizons. In bulk soil with the organic horizon, the activities of NAGase and glucosidase were slightly higher and the activity of xylosidase was lower than those in bulk soil without the organic horizon (Fig. 4). However, the root system, organic horizon, and interaction between the root system and organic horizon had no significant effects on the extracellular enzyme activities in bulk soil, except for on xylosidase activity, which was significantly higher when root systems were present (Fig. 4).

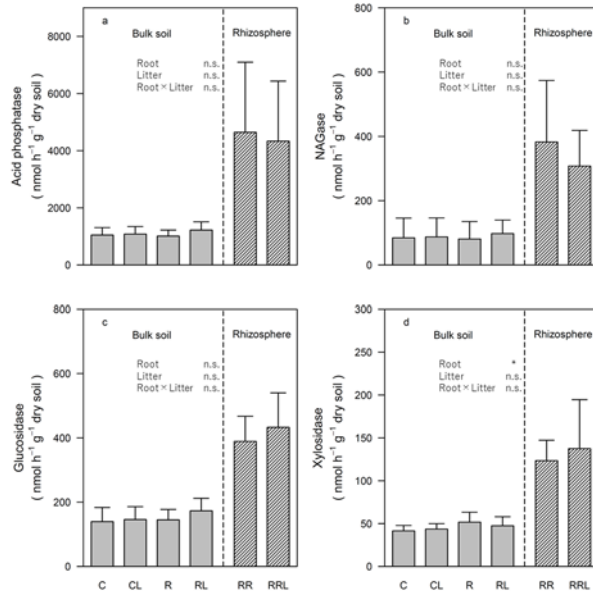


Fig. 4 Activities of (a) acid phosphatase, (b) NAGase, (c) glucosidase, and (d) xylosidase in the bulk soil (gray bars) and rhizosphere soil (shaded bars). The bars and error bars represent the means and standard deviations, respectively ($n = 4$). C, CL, R, and RL represent each treatment. RR and RRL represent the rhizospheres of R and RL, respectively. Root, Litter, and Root \times Litter in the upper center of each figure represents the results of the two-way ANOVAs with a random effect for the enzyme activities in bulk soils. The symbols are as follows: n.s.: $p \geq 0.05$, ***: $p < 0.001$

Discussion

Plant roots directly affect soil N availability in two ways, that is, through N uptake and through their influence on the decomposition and N mineralization processes via the priming effect. Although the fine roots of *Q. crispula* did not significantly alter the N pools or N leaching in this study, this does

not suggest that the fine roots had no effect on N availability; rather, the fine roots accelerated these processes and took up the excess N mineralized by this acceleration effect. Thus, in this study, we showed that this is the mechanism by which the fine roots of canopy *Q. crispula* affect N availability in forest soils by using our novel modified *in-situ* resin core method, which accounts for both N uptake and priming by fine roots.

Our results did not contradict hypothesis (1), that is, extracellular enzyme activities were stimulated by fine roots, and remarkably so in the rhizosphere (Fig. 4). This was consistent with the findings of previous studies in both herbaceous and woody species (Phillips and Fahey, 2006; Zhu et al., 2014; Kumar et al., 2016). Meier et al. (2017) reported that labile organic C inputs, including root exudates, enhanced the rapid cycling of organic N and the rapid turnover of organic N-containing compounds, such as chitin, which is the substrate for NAGase, and increased N availability in the soil. The positive effect of roots on extracellular enzyme activities, especially xylosidase, was also observed in bulk soil in this study, although NAG activity was slightly reduced by the presence of fine roots in the absence of an organic horizon (Fig. 4). *Quercus crispula* has symbiotic relationships with ectomycorrhizal fungi (Obase et al., 2007). Previous studies have shown that ectomycorrhizal fungi extensively extend their mycelium into the bulk soil (Cairney, 2012; Soudzilovskaia et al., 2015), expanding the effect of plant-derived C into soil that is distant from the roots (Meier et al., 2015; Gorka et al., 2019). Xylo-oligomers, the substrate for xylosidase, are strong inhibitors of the

enzymatic degradation of recalcitrant organic matter (Qing et al., 2010). Thus, stimulation of xylosidase activity would enhance recalcitrant organic matter decomposition. In other words, at least a part of the root-distant soil area regarded as bulk (non-rhizosphere) soil in this study would be the hyphosphere, and the fine roots of *Q. crispula* might accelerate the decomposition of recalcitrant SOM in the bulk soil via the mycelium of their symbiotic mycorrhiza. Therefore, our results indicate that the fine roots of *Q. crispula* stimulated the decomposition of recalcitrant SOM within the rhizosphere and bulk soil, and enhanced the rates of N turnover and cycling, especially in the rhizosphere. As an increase in SOM decomposition could increase N mineralization (Dijkstra et al., 2009), the fine roots of *Q. crispula* might enhance N mineralization via rhizosphere and hyphosphere priming effects. However, the excess inorganic N mineralized by the stimulatory effects of the roots would then be readily taken up by the fine roots in the cores, as discussed below.

As for hypotheses (2) and (3), the presence of fine roots did not significantly affect either the size of the N pool or the amount of inorganic N that was leached (Figs. 2 and 3), regardless of their potential to stimulate SOM decomposition via the priming effect, suggesting that N_{uptake} could be almost equal to N_{induced} in this study (Table 4). In temperate forest soil, root-accelerated mineralization can account for up to one-third of the total N mineralization (Finzi et al., 2015). However, the magnitude and direction of the effect of plant roots on N availability and N loss by leaching are determined by the balance between plant N uptake and priming. Dijkstra et al. (2009)

showed that the seedlings of woody species decreased the inorganic N pool by utilizing it along with mineralized N; however, they observed a significant positive RPE on SOM decomposition. Herman et al. (2006) reported that gross mineralization was approximately 10-times higher in rhizosphere soil than in bulk soil, but N uptake by slender oats exceeded the amount of N supplied by N mineralization in the rhizosphere. In this study, the lower density of fine roots in the incubation pipes suggested that the amount of N taken up by the fine roots was lower than that under field conditions. The “actual” uptake of N by fine roots also could differ from the “potential” estimated as N_{uptake} because N availability might affect N uptake (Moreau et al., 2015; Liao et al., 2018). These are limitations of this study. Therefore, further research is required to determine whether canopy trees increase N retention in forest ecosystems. However, even if root-induced changes in N mineralization are quantitatively important, our results suggest that the fine roots of canopy trees do not increase N loss through leaching from the forest ecosystem because almost all of the N mineralized by root-accelerated mineralization was immediately taken up by the trees.

Furthermore, when the organic horizon was present, N_{induced} was almost zero (Table 4). There are two possible explanations. One is that there was a lower RPE on N dynamics with higher nutrient availability. The magnitudes of the RPEs on extracellular enzyme activities related to N and phosphate acquisition in cores containing the organic horizon were slightly lower (Fig. 4). As suggested by the differences in N leaching between C and CL (Fig. 3), some amount of substrates and

energy sources for microbes (including organic N) would be supplied by the organic horizon material added to the cores; these supplements might lead to a reduced RPE. Based on the microbial nutrient mining theory of RPE, root exudates are considered to stimulate SOM decomposition more in nutrient-limited environments than in nutrient-rich ones (Craine et al., 2007; Fontaine et al., 2011; Dijkstra et al., 2013; Cheng et al., 2014). For example, Phillips and Fahey (2008) reported that N fertilization reduced the RPEs on microbial biomass, extracellular enzyme activity, and the net mineralization rate. Moreover, although it is still unclear whether canopy trees increase root exudation under nutrient limitation (e.g., Phillips et al. 2008, 2011; Meier et al. 2020), plants allocate more C to the belowground parts under nutrient-limited conditions (Hill et al., 2007; Chowdhury et al., 2014). The magnitude of the RPE is positively correlated with the quantity of labile C from fine roots (Yin et al., 2013; Shahzad et al., 2015). Studies have suggested that a higher RPE results when nutrient availability for plants is higher (Dijkstra et al., 2009; Bengtson et al., 2012); however, the release of organic C is a significant C cost for plants (Uren, 2007). Although significant amounts of root exudates have been observed under natural condition in this site (Nakayama and Tatenno, 2018), it is still possible that canopy trees decrease the priming effect on N mineralization by decreasing labile organic C release when N limitation is less severe.

The second possible explanation for why N_{induced} is almost zero in treatments with an organic horizon is that the difference between “actual” and “potential” N uptake, which were derived from the

inappropriate assumption of our estimation. Moreau et al. (2015) investigated 11 crops and model species and found that higher soil N availability led to higher N uptake per gram of fine roots, although the extent of this effect varied by plant species. Maeda et al. (2018) reported that the combination of soil N mineralization and belowground net primary production affected tree nitrogen uptake, while soil N mineralization alone did not significantly correlate to tree N uptake in a natural deciduous-broad leaved forest in East Hokkaido, the same region as that in this study. Indeed, the RPEs on the activities of acid phosphatase and NAGase were slightly lower in the pipes with organic horizons than in those without organic horizons, but the difference was not statistically significant. Therefore, there is a possibility that the actual N uptake might be higher in the pipes with organic horizons than in those without organic horizons because of nutrient availability, and the actual root-induced N increments might be similar regardless of the difference in the treatment of the organic horizon.

In northern temperate forests, nitrogen is often regarded as the limiting factor for forest primary production (Vitousek and Howarth, 1991; LeBauer and Treseder, 2008). At the same time, the soil in the study site is an Andosol, which tend to have high total phosphorus (P) content, mainly as organic P, but low available P (Borie and Rubio, 2003). Inorganic P deficiency increases the release of phosphatase by plants and the activity of phosphatase (Almeida et al., 1999; Oberson et al., 2001). This would explain why acid phosphatase activity was higher than the activities of the other enzymes observed in this study. Therefore, our extracellular enzyme activity and N budget results might be

specific to Andosols. Further research using the modified resin-core method in other soil types will provide detailed insights into the relationships between canopy tree fine roots and N dynamics.

In this study, we modified the *in-situ* resin core method to investigate the effect of the fine roots of a specific tree species on the N cycle in forest soil. The results of this study suggest that the fine roots of *Q. crispula* stimulated SOM decomposition and N mineralization; however, they did not increase either the inorganic N pool size or N loss from the ecosystem because canopy trees took up almost all inorganic N mineralized by the priming effect of fine roots. Nevertheless, this study was limited by the fact that fine roots used in the cores were less dense than those under natural conditions, and the possibility that “actual” root N uptake differs from the “potential” estimated N_{uptake} . Furthermore, we only investigated the effect of roots of a single species and neglected the effects of interspecific interactions, whereas a natural forest ecosystem contains the roots of various plant species. However, the method used in this study is effective for investigating the relationships between canopy trees and forest soil N cycling, while considering both N uptake and priming. Future research using this method with higher root densities and with the fine roots of two or more tree species will provide more insights into the interactions between the fine roots of canopy trees and N cycling in forest soil.

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767 Table 1 Total C and N contents and C/N ratios of the mineral soil and the organic horizon used in the incubation.

	Total C (%)	Total N (%)	C/N ratio	Dry weight used in the incubation (g pipe ⁻¹)
Organic horizon				
Relatively fresh	48.94 ± 0.75	1.10 ± 0.14	45.0 ± 5.8	0.25
Partially decomposed	45.71 ± 1.96	1.26 ± 0.21	37.4 ± 7.3	2.46
Passed through 9 mm mesh	42.75 ± 3.16	1.67 ± 0.11	25.6 ± 1.2	1.25
Mineral soil				
	8.58 ± 0.99	0.59 ± 0.07	14.6 ± 0.3	129.64

768 Note: Values represent the mean ± standard deviation.

769 Table 2 Equations used to estimate nitrogen uptake, nitrogen mineralization, nitrification, and the priming effect on nitrogen mineralization.

Parameter	Abbreviation	Equation
Nitrogen uptake		
Nitrogen demand of <i>Q. crispula</i>	N _{demand}	Sum of nitrogen contents in each part of net primary production
Potential nitrogen uptake per dry weight of fine roots under natural conditions	Nat N _{uptake}	N _{demand} / Fine root density of <i>Q. crispula</i> (0–50cm) in study plots
Potential nitrogen uptake by fine roots in incubation pipes	N _{uptake}	Nat N _{uptake} × Fine root densities in incubation pipes
Nitrogen dynamics		
Net nitrate increment	Nit _{inc}	Annual leaching + increment of pools of NO ₃ ⁻ -N
Net inorganic nitrogen increment	N _{inc}	Annual leaching + increment of pools of NO ₃ ⁻ -N and NH ₄ ⁺ -N
Fine root induced increment of inorganic nitrogen	N _{induced}	N _{min,R} – N _{min,C} + N _{uptake}

770 Note: N_{min,R} and N_{min,C} represent N_{min} of samples with and without *Q. crispula* fine roots, respectively.

771

772

773 Table 3 Production, N concentration, and N demand for *Q. crispula* overall and for individual plant parts.

	Production (g/m ² /year)	N concentrations (%)	Estimated N demand (g-N/m ² /year)
Total NPP	601.5 ± 76.5		6.90 ± 1.01
Leaf litter	256.7 ± 36.1	1.82 ± 0.08	4.67 ± 0.66
Other litter	21.2 ± 9.14	1.46	0.31 ± 0.13
Branch	96.3 ± 11.7 ^{*1}	0.58 ^{*2}	0.56 ± 0.07
Trunk	152.2 ± 23.0 ^{*1}	0.39 ^{*2}	0.59 ± 0.09
Coarse root	62.2 ± 10.2 ^{*1}	0.45 ^{*2}	0.28 ± 0.05
Fine root (0-10cm)	18.3 ± 23.5	1.43 ± 0.16	0.26 ± 0.34
Fine root (0-50cm)	34.1 ± 43.8 ^{*3}		0.49 ± 0.63 ^{*3}

774 Note: NPP represents the net primary production. Values represent the mean ± standard deviation (n = 4 for the production of branch, trunk, and coarse roots,
775 and the N concentration in fine roots; n = 8 for leaf litter; n = 32 for fine root production).

776 *1: Branch, trunk, and coarse root productions were calculated based on the allometric relationship (Takagi et al., 2010) and increase in diameters at breast
777 height.

778 *2: N concentrations in the branch, trunk, and coarse root were based on a previous study (Takagi et al., 2010).

779 *3: Fine-root production (0–50 cm) was estimated by the vertical distribution of the fine roots of *Q. crispula* in the adjacent plot (Tateno et al., 2020).

780

781 Table 4 Net N dynamics, estimated potential root N uptake based on the root density in the modified resin core, and the root-induced annual inorganic N

782 increment (0–10 cm soil depth).

	With organic horizon		Without organic horizon	
	With root	Without root	With root	Without root
Net N dynamics				
Nit _{inc} (g-N m ⁻² year ⁻¹)	6.61 ± 1.04	7.14 ± 1.09	4.93 ± 0.90	4.99 ± 1.13
N _{inc} (g-N m ⁻² year ⁻¹)	6.53 ± 1.11	7.01 ± 1.12	4.78 ± 0.87	4.92 ± 1.11
N _{uptake} (g-N m ⁻² year ⁻¹)	0.52 ± 0.18		0.53 ± 0.13	
N _{induced} (g-N m ⁻² year ⁻¹)	0.04 ± 0.55		0.32 ± 0.48	

783 Note: N_{nit}, N_{min}, N_{uptake}, and N_{primed} represent the annual nitrification, annual nitrogen mineralization, annual nitrogen uptake, and annual primed nitrogen

784 mineralization by the fine roots in the incubation pipes, respectively. The values represent mean ± standard deviation.

785