



## Fine Root Size and Morphology of Associated Hyphae Reflect the Phosphorus Nutrition Strategies of European Beech Forests

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Roots are among the major controls of nutrient and C cycles and together with mycorrhizal fungi they are assumed to play a key role especially in the P nutrition of forest ecosystems. Current publications emphasized that the size distribution of fine roots reflects the crucial impact of roots on biogeochemical cycles. However, we know hardly anything about the spatial distribution of fine root size classes and their specific surface as well as the distribution of mycorrhizal fungi among different size classes in undisturbed soils. We used a novel method based on epifluorescence microscopy to analyze fine roots in undisturbed soil samples. We expected that based on these analyses proposed differences between P-rich and P-poor soils get clearer than based on routine methods for fine root analysis. This was examined at two European beech forests in Germany at silicate rock that differ in P supply by the parent material. We analyzed the fine root frequency, size distribution and surface based on resin impregnated cross sections taken from undisturbed soil samples. Fine roots were classified according to their size and morphology of associated ectomycorrhiza (ECM). After staining with acridine orange these root traits were analyzed and quantified with epifluorescence microscopy. We found that more than 82 % of the absorbing surface was associated with roots having a diameter smaller than 100  $\mu$ m. The fine root surface area present per square meter of soil was 388  $m^2$  at the P-poor site and 220  $m^2$  at the P-rich site. In addition, the percentage of mycorrhization of the fine root surface was 47 % at the P-poor site and only 38 % at the P-rich site. The biggest root length density and the highest absorbing fine root surface with mycorrhization and abundance of extramatrical mycelia (EM) was observed in the forest floor (+4 to 0 cm) at the P-poor site and in the subsoil (-79 to -100 cm) at the P-rich site. Our results confirm that beech trees adapt their root traits according to P availability showing higher absorbing surface at the P-poor site compared to the

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P-rich site. In contrast, at the soil-profile scale, rooting density (RLD) and mycorrhization increased with P availability thereby showing an efficient way of root and mycorrhizal fungi placement. Overall, distinct differences in fine root traits, between the P-rich and P-poor site were most evident for fine roots smaller than 100  $\mu$ m.

Keywords: phosphorus-nutrition, fine root length, ectomycorrhiza, beech forests, root size class, epifluorescence microscopy, polished soil sections

## **1. INTRODUCTION**

Roots are less accessible for researchers than above-ground plant compartments and are widely assumed to bear still a lot of unidentified possibilities of adaptation (Finér et al., 2011). Current papers indicate the prominent role of roots in C sequestration (Ruess et al., 2003; Clemmensen et al., 2013) and nutrient cycling (Johnson and Turner, 2019). The high relevance of roots for phosphorus nutrition of higher plants originates from the low mobility of P in soils (Raghothama and Karthikeyan, 2005). At many sites, P is among the limiting factors of forest growth (Lang et al., 2017). It has been shown that beech forests evolved many mechanisms to adapt to P scarcity. Since declining P leaf contents observed for European beech forests (Jonard et al., 2015; Talkner et al., 2015) might indicate disturbance of those mechanisms they are currently in the center of scientific debate. Lang et al. (2017) found a shift in P nutrition strategies of beech forest ecosystems, depending on the P supply provided by the soils. At sites with low stocks in soil P, beech forests maintain their P supply by recycling of P from organic sources, while the acquisition of P from mineral resources controls P nutrition at high soil P stocks (Raghothama and Karthikeyan, 2005). Till now, it is unknown, if fine root spatial and size distribution as well as mycorrhization reflect nutrition strategies (Lopez-Bucio et al., 2000; Plassard and Dell, 2010; Hauenstein et al., 2018). In general, size distribution and surface area of very small roots (<100  $\mu$ m) and the morphology of the associated mycorrhizal fungi have hardly been analyzed. This is especially true for samples with undisturbed soil structure. About 95% of all trees in forest ecosystems, especially in boreal and temperate biomes, have ectomycorrhiza symbiosis (Smith and Read, 2009). In terms of P uptake, the extramatrical mycelium (EM) produced by ectomycorrhizal fungi (ECM) is a key factor (Read, 1984; Plassard and Dell, 2010). It improves P uptake by increasing the size of the explored soil volume (Lynch, 1995) whereby the depletion zone may extend far beyond the rhizosphere (Bolan, 1991; Marschner and Dell, 1994; Tibbett and Sanders, 2002) and thus overcomes the limits of the slow diffusion of P in the soil, which is assumed to be not more than a few millimeters (Williamson and Alexander, 1975; Schachtman et al., 1998; Plassard and Dell, 2010; Demand et al., 2017). Furthermore, Nylund and Wallander (1992) determined that the production of the EM for Pinus sylvestris L. is induced by P starvation. Thus, we assume that European beech trees of the above-mentioned recycling and acquiring systems show different rooting density, depth distribution, and mycorrhization depending on P availability.

Most published root data is based on mass or length of washed-out roots (Böhm, 1979; Colpaert et al., 1992; Van Noordwijk et al., 2000). However, this procedure has problems regarding the size distribution of roots in soils. It has been found that especially fine roots (<1 mm) are underestimated (Bengough et al., 1992; Rousseau et al., 1994). The quantification of hyphae spreading out from the Hartig net to build extramatrical mycelium is even more challenging (Colpaert et al., 1992; Rousseau et al., 1994). A variety of methods have been developed to quantify the biomass of extramatrical mycelium in soils, including mini-rhizotrons and in-growth mesh bags. Yet, due to disturbance of soil samples or artificial growing conditions, these methods are inadequate to quantitatively describe roots and their mycorrhizal traits in soils. Consequently, to provide information on the effective absorbing root surface area, multiple studies have requested that roots should be characterized and quantified in undisturbed soil samples (Jones et al., 1990; Rousseau et al., 1994; Hinsinger et al., 2011; Peñuelas et al., 2013). We suggest a micro morphological approach to quantify and characterize fine roots which has also been used by Tippkötter et al. (1986), Altemüller (1989), and Schack-Kirchner et al. (2000). This means, shortly, that roots are left in their soil environment but are analyzed on two-dimensional areas such as thin sections or faces prepared otherwise. Our study aim was to analyze those characteristics of fine roots (<2 mm) with major relevance for P uptake: size distribution, surface area, specific root length, and the mycorrhization as well as the morphology of mycorrhizal fungi within undisturbed soil environment. We compared soils from two European beech forests, one developed from Prich substrate, the other from P-poor substrate. We expected that based on micro morphological root analyzes proposed differences between P-rich and P-poor soils get more clear than based on routine methods for fine root analysis. Our approach was based on epifluorescence microscopy of resin impregnated polished soil sections taken from undisturbed soil cores. To our knowledge, this study is the first approach of in situ quantification of the absorbing root surfaces in their undisturbed environment.

## 2. MATERIALS AND METHODS

## **Site Characteristics**

Soil and stand characteristics including P stocks of the study sites Bad Brueckenau (BBR, P-rich site) and Unterluess (LUE, P-poor site) are reported in **Table 1**. As a proxy for P availability, citric acid extractable  $P_{inorganic}$  according to Manghabati et al. (2018) is listed per soil horizon in **Table 2**. The experiment is focused

Properties	BBR	LUE	
Geographical coordinates	50°35′N, 9°92′E	52°83′N, 10°36′E	
Soil type (WRB, 2015)	Dystric skeletic cambisol (Hyperhumic. Loamic)	Hyperdystric folic cambisol (Arenic. Loamic. Nechic. Protospodic)	
Parent material	Basaltic rock	Sandy till	
Humus form (Ad-hoc-AG Boden, 2005)	Mull-like Moder	Mor-like Moder	
Texture—topsoil (WRB, 2015)	Silty clay loam	Loamy sand	
Texture—subsoil (WRB, 2015)	Loam	Sand	
Elevation [m]	809	115	
Precipitation [mm]	1,031	779	
Total basal area m <sup>2</sup> ha <sup>-1</sup>	35.6 Fagus sylvatica (99%), Acer pseudoplatanus (1%)	36.7 Fagus sylvatica (91%), Quercus petraea (9%)	
Stand age [a]	137	132	
Stem density [n ha <sup>-1</sup> ]	595	520	
P Stocks to 1 m [kg ha <sup>-1</sup> ]	9,040	164	

TABLE 1   Site and stand characteristics of the study sites Bad Brueckenau
(BBR) and Unterluess (LUE) (Lang et al., 2017).

exclusively on four soil horizons per site (Oa, Ah/AhE, Bw/Bsw, and Cw). BBR, is located in the Rhoen mountains in North Bavaria, and LUE, is in the North German lowlands. Both study sites belong to the Level II intensive monitoring plots of the Pan-European International Cooperative Program on assessment and monitoring of air pollution effects for the forests (ICP Forests) under the supervision of UNECE (Lorenz, 1995). Though the stocks and availability of P substantially differ between the two sites, leaf concentrations of P are similar (Lang et al., 2017).

## 2.1. Root-Mycorrhiza Investigations

2.1.1. Root System Sampling and Soil Preparation For the quantitative investigation of fine roots (<2 mm), we used a three-level hierarchical sampling design (Figure 1). On the first level, we sampled undisturbed soil cores at three sampling points (three beech trees) considered to be representative of the sites BBR and LUE and with low coverage of herbs or shrubs to avoid non-beech roots and at a distance from the next stem of about 1 m. Soil cores with a diameter of 8 cm and a length of 20 cm were taken with a root auger up to 1 m in depth. In BBR, we had to take clods below 40 cm instead of soil cores due to the high stone content. Immediately after sampling, we identified and counted the visible fine roots (1-2 mm) (Kreutzer, 1961) by the core break method (Böhm, 1979). On the second level, we subsampled the surrounding of four randomly chosen visible fine roots (1-2 mm) per analyzed soil horizon with "mini cores" for microscopic analysis.

These mini cores (Figure 1) are tubes formed out of perforated aluminum plate with a diameter of 1.5 cm and a

length of 2 cm. These were pushed into the soil around a fine root (1–2 mm). The mini cores and the Cw clods were next placed in rectangle casting molds (EPDM rectangle mold,  $55 \times 30 \times 22$  mm, Buehler, Esslingen am Neckar, Germany).

For the third level, immediately after cutting out and placing the mini cores were fixed in 2.5% glutardialdehyde aqueous solution (Tippkötter et al., 1986) to stop microbial activity and to conserve roots and other biological tissues. Dehydration was then performed using a five-step ascending acetone concentration of 50%, 80%, and three times 100% for 3 days to prevent shrinking or deformation (Tippkötter and Ritz, 1996). The dehydrated soil samples were then embedded in a low viscosity epoxy resin (Serva Electrophoresis GmbH; Heidelberg; Germany, Application note "Modified Spurr," "Modification E") (Serva Electrophoresis GmbH, 2007). After complete filling of the pore space, acetone was replaced by resin. The resin content was sequentially increased from 20 to 100% in five steps based on preliminary tests. To ensure that all pores were filled with epoxy resin, the actual impregnation was carried out with 100% epoxy resin under a vacuum for 24 h.

The final polymerization took place for 16 h at 70°C. The cured soil cores were mounted onto aluminum plates with hot melting adhesive and were cut horizontally into three to five 1–2 mm slices using a diamond saw. These slices were ground with sand paper in declining grid size from P320 to P800 and P1200, and finally polished with  $Al_2O_3$  powder with declining grain size from 5, 1, and 0.3  $\mu$ m. Low viscosity erosion oil (EDM180 from Ilocut, Diermeier Energie GmbH, Munich, Germany) was used as cutting and grinding fluid to avoid swelling (Schack-Kirchner et al., 2000). The individual soil samples were finally glued onto the microscope slides with a two compound epoxy adhesive glue and cleaned with paraldehyde. Since the soil slices did not perfectly correspond to the mini core surface area, we quantified the soil filled surface area of each soil slice with image analysis (ImageJ, Version1.51f).

The surface of the polished soil slices was stained with acridine orange. According to Schack-Kirchner et al. (2000) for the coloring of hyphae embedded in polyester resin was modified as follows: In preliminary tests we found an optimum staining contrast in epoxy resin specimen by using a mixture of seven parts 0.1% acridine orange in 100% ethanol to three parts 0.1% acridine orange in 10% hydrochloric acid. It was applied for 10 min and the staining was done in closed boxes to prevent the evaporation of ethanol. After the dye was removed with a cloth, the soil slabs were left to stand for a further 20 min before being examined by Zeiss Axio Zoom V16 fluorescence zoom microscope (Carl Zeiss AG; Jena, Germany). For the fluorescence illumination, a HXP 200C illuminator with a mercury short-arc reflector lamp was used. The microscope was equipped with a PlanNeoFluar Z2.3x/0.57 FWD 10.6 mm objective and a filter combination of BP450-490/FT510/LP515.

The digital microscope camera AxioCam HRC (Carl Zeiss AG; Jena, Germany) has a CCD sensor type with a pixel resolution of 1.4 megapixel with 1,388  $\times$  1,040 with a pixel size of 6.45  $\mu$ m. The pictures were further analyzed with the imaging software ZEN 2012 (Version 1.1.2.0, Carl Zeiss AG; Jena, Germany).

Site	Horizon	Soil depth	рН <sub>Н2</sub> о	Pcitr	С	N	CEC	BS
		cm		$mg kg^{-1}$	%	%	meq kg $^{-1}$	%
BBR	Oa	+2 to 0	5.3	326	48.2	2.07	543	79
BBR	Ah	0 to -7	4.2	39	9.7	0.65	261	10
BBR	Bw	-33 to -51	4.9	64	3.4	0.25	149	26
BBR	Cw	-79 to -100	5.7	141	1.1	0.09	148	45
LUE	Oa	+4 to 0	3.3	54	46.3	1.75	557	13
LUE	AhE	0 to -7	3.5	10	9.7	0.38	109	8
LUE	Bsw	-65 to -79	4.6	3	0.4	0.03	38	7
LUE	Cw	-88 to -100	4.6	3	0.2	0.02	23	9

TABLE 2 | The citric acid extractable P for the respective sampled soil horizon (Oh, Ah, Bw, and Cw) of the two study sites Bad Brückenau (BBR) and Unterluess (LUE) (Lang et al., 2017).



By illumination with blue light, fresh plant material, hyphae, and bacteria fluoresced in a spectrum from yellow to yellowgreen. However, a color difference between hyphae and other plant material was not visible. Therefore, roots and fungi were classified and determined by differences in morphology that could be easily distinguished such as diameter, form, or the appearance of hyphae mantle. The respective cell structures of roots and fungi could be clearly distinguished.

A total of 384 microscopic samples distributed evenly over both sites were evaluated. We included only intact roots in our data set. First, we differentiated between fine roots having a diameter of 1-2 mm, and those having a diameter <1 mm in



ZEN 2012 (Figure 2). We used a spherical graphic primitive interactively placed into the boundaries of the root cross section to measure the minimum ferret diameter representing the diameter even when roots are cut non-perpendicular to their axis. Second, for each root, the existence of mycorrhization was analyzed based on presence/absence of hyphae mantle. Since melanine hyphae are not stained by acridine orange, whereas hyaline hyphae are, we were able to differentiate between melanine and hyaline hyphae as described in Schack-Kirchner et al. (2000) (Figure 2).

#### 2.1.2. Classification of Fine Roots

Roots were assigned to different morphological classes according to diameter, existence of a hyphae mantle, type of hyphae, and emanating hyphae as an indication of EM (**Figure 2**). For this classification, each root cross section on the specimen was assessed. With our microscopic method it is possible to differentiate low order absorptive roots with no secondary development and roots with mainly transport functions based on absence/presence of fully developed cortexes and xylem as suggested by McCormack et al. (2015). Nearly none of our cross sections showed fully developed cortex.

Microscopic images of different fine root classes are shown exemplary in **Figure 3**.

#### 2.1.3. Calculations

For the first level (**Figure 1**) the fine root (1–2 mm) frequency per surface area was calculated as:

$$\frac{N_{1-2 mm}}{A_{Ac}} \tag{1}$$

with  $A_{Ac}$  as the area of the vertical root-auger core surface and  $N_{1-2 mm}$  as the number of visible fine roots tips (1–2 mm). For the second level (**Figure 1**) the fine root (<1 mm) of the mini core cross section were counted to calculate the frequency:

$$\frac{N_{<1\ mm}}{A_{Mc}}\tag{2}$$

With  $A_{MC}$  as the area of mini core surface and  $N_{<1 mm}$  as the number of fine roots (<1 mm). Finally, the number of fine roots (<1 mm) was averaged across the vertical repetitions of up to five slices. By assuming that the detected cross sections of roots were part of an isotropic multitude of vectors in 3 D space, fine root length densities (RLD/cm cm<sup>-3</sup>;  $\frac{L_R}{V_S}$  in Weibel, 1980) were derived from the number of cross sections per area (Weibel, 1980; Schack-Kirchner et al., 2000; Van Noordwijk et al., 2000):

$$RLD = \frac{L_R}{V_S} = \alpha * \frac{N_R}{A_S}$$
(3)



hyaline hyphae mantle and hyaline extramatrical mycelium. Further information can be found in Figure S1.

where  $L_R$  is the root length [cm],  $V_S$  is the soil volume [cm<sup>3</sup>],  $\alpha$  is the dimensionless factor of isotropy (we assumed complete isotropy with  $\alpha = 2$ ; Weibel, 1980),  $N_R$  is the number of the counted fine roots, and  $A_S$  is the soil area. Based on the specific RLD ( $L_R/V_S$ ), thickness of each horizon H and average diameter  $\overline{d}$  for each fine root trait per horizon, we calculated the fine root surface area index (RAI/m<sup>2</sup> m<sup>-3</sup>) per soil volume:

$$RAI = \pi * \bar{d} * RLD * H \tag{4}$$

The RAI was related to the stand area, separately for each genetic soil horizon, as a proxy for the absorbing fine root surface

considering the thickness of each horizon. The calculations were carried out for the different fine root classes.

#### 2.1.4. Statistics

From each mini core we assessed four to five cross sections. The root-observations have been averaged because these samples are not statistically independent. From each soil horizon we obtained 12 averaged mine core data sets of root counts originating from three soil cores. Because variability of root counts within soil cores was much lower than between soil horizons and average results of the respective soil cores individually reflected already the overall horizon pattern, we regarded the averaged mine core results as statistically independent, so we have



N = 12. To test the observed differences in medians for statistical significance we used the multiple non-parametric multiple Dunn's test with Bonferroni correction (Kruskal-Wallis-test, p < 0.05) (https://cran.r-project.org/web/packages/dunn.test/dunn.test.pdf).

## **3. RESULTS**

In total, we measured and classified 6,969 fine roots (<1 mm) in BBR and 13,587 fine roots (<1 mm) in LUE. Diameters >500  $\mu$ m were observed rarely, the frequency maximum was



usually below 100  $\mu$ m. (**Figure 4**). Roots (d > 1 mm) were found only sporadically. By far the highest number of fine roots were found in the topsoil of LUE.

In general, the degree of mycorrhization was lower at BBR than at LUE. The least share of mycorrhizal roots

at BBR was observed in the Oa horizon (<9.7%) and at LUE in the Cw horizon (33.1%, **Figure 5**). In comparison, horizons with the highest portion of mycorrhizal roots were the Cw horizon in BBR (41.5%) and in the Bsw horizon in LUE(>51%).

**TABLE 3** | The calculated root length density (RLD) and the ratio of root length density (RLD) <1 mm to root length density (RLD) (1–2 mm) are compared to the RLD<sub>sieving-scanningmethod</sub> of sieving-scanning method of Lang et al. (2017) at the same sites.

Site	Horizon	RLD <sub>1-2 mm</sub> cm cm <sup>-3</sup>	RLD <sub>&lt;1 mm</sub> cm cm <sup>-3</sup>	$\frac{R_{<1 mm}}{R_{1-2 mm}}$	RLD <sub>sieving-method</sub>
BBR	Oa	11.5	120	10	0.29
BBR	Ah	4.75	31	7	2.01
BBR	Bw	1.7	20	12	0.46
BBR	Cw	0.5	54	106	0.12
LUE	Oa	8.08	191	24	5.37
LUE	AhE	4.2	164	37	1.38
LUE	Bsw	0.46	41	89	0.27
LUE	Cw	0.05	33	60	0

The mean root length density and standard deviation for all mini core samples per soil core horizon and site are listed in **Figure S4**. The mean root length density. Furthermore, the box plots of the RLD over all mini core mean values together with the result of the Dunn-test for each median are represented in **Figure S5**.

The highest portion of fine roots (<1 mm) with extramatrical mycelium was found in the Cw horizon of BBR (13%) and in the Bsw horizon of LUE (16%). In addition, at BBR the portion of hyaline mycelia was higher than at LUE.

The highest relative portion of fine roots (<1 mm) with EM were found in the Cw horizon of BBR (12.6%) and in the Bsw horizon of LUE (16.1%), the lowest portion were in the Oa horizon of BBR (0.3%)and in the AhE and the Bsw horizon of LUE (3.2%). Melanine mycorrhization dominated in the topsoil of LUE and in the Cw horizon of BBR, hyaline mycorrhization dominated in the Ah horizon of BBR and in the Bsw horizon of LUE. In the Oa, AhE, and Cw horizons of LUE and in the Bw and Cw of BBR, the EM was dominated by melanine hyphae, whereas the other horizons of BBR and LUE were dominated by hyaline EM.

In BBR, the RLD<sub>1-2 mm</sub> [cm cm<sup>-3</sup>] consistently decreased with depth while the RLD<sub><1 mm</sub> [cm cm<sup>-3</sup>] decreased from the Oa to the Bw horizon but rose again in the Cw horizon (**Table 3**). In LUE, both fine root length densities revealed a steady decrease with depth (**Table 3**). At both sites, the ratio of RLD<sub><1 mm</sub> to RLD<sub>1-2 mm</sub> increased with depth except the Ah horizon in BBR. However, in LUE this ratio was consistently two to six times higher than in BBR.

From each soil horizon we obtained 12 averaged mine core data sets of root counts originating from 3 soil cores. All soil cores are similar (**Figure S4**) which is why all mini core samples can be considered as an independent sample (**Figure S4**). For these we calculated root length density (RLD) and the ratio of root length density (RLD) <1 mm to root length density (RLD) (1–2 mm) are represented. Furthermore, we compared the RLD of our method with the RLD *sieving*-*scanningmethod* of a routine sieving-scanning method of Lang et al. (2017) also performed at both sites (**Table 3**).

In the following it is analyzed whether the horizons within the profiles or whether the same horizons of the different profiles

differ (**Figure 6**). It can be clearly seen that the variability is between the soil cores, which means that the heterogeneity within the horizons is not so significant.

Based on the RLD and mean diameter (**Figure 5**), we calculated a root surface area index (RAI)  $[m^2 m^{-3}]$ , (**Figure 7**). If RAI is related to the genetic horizon depths up to 1 m soil depth, the weighted RAI is obtained (right bar). We regard this weighted cumulative RAI as a measure of as the potentially absorbing area per stand area. In LUE the total this RAI is almost twice as high and the mycorrhizal RAI almost three times as high as in BBR. The horizon-specific RAIs reveal the disproportionately high contribution of the only 11 cm thick organic topsoils (Oa and AhE) in LUE and the only 2 cm thick Oa horizon and 21 cm Cw horizon in BBR.

In **Figure 8**, we have related the RLD to the soil concentration of available P, which reaches 300  $mg kg^{-1}$  in the Oa of BBR and only 50  $mg kg^{-1}$  in the Oa in LUE. The general pattern at both sites was an increase in total RLD of fine roots <1 mm with P availability regardless of soil depth. However, the total mycorrhizal root length and its share in the total root length do not show a clear pattern. To test the influence of P on the RLD distribution, we tested correlations between the different RLDs and P, soil texture, pH, base cations, etc. for each site separately (**Figures S2, S3**). However, we could not find any influence on the root length density in our correlations, except the soil P content (**Figures S2, S3**).

## 4. DISCUSSION

## 4.1. The Method Applied

We observed that 82% of the absorbing root surface was bound to fine roots having a diameter smaller than 100  $\mu$ m. These results were in contrast to results reported by Weemstra et al. (2017) that most of the fine roots of European beech were part of the diameter classes 200–750  $\mu$ m. In agreement, also other studies indicated low abundance of root diameter classes smaller than 500  $\mu$ m (e.g., Leuschner et al., 2001). A direct comparison of visual and microscopic scanning approaches provided in Table 3 supports that visual methods in particular fail to quantify the roots smaller than 1 mm (Table 3). In agreement, also the roots surface seems to be strongly underestimated by conventional methods (Jackson et al., 1996; Kirfel et al., 2019) sieving-scanning or washing-scanning cause losses especially of small fine roots. Furthermore, picking out roots manually, also results in an underestimation of fine roots since only roots seen by eyes without the support of a microscope, are selected.

Functional traits of roots are in the center of current research (McCormack et al., 2015). An advantage of our method is the possibility to differentiate between absorptive roots and roots with mainly transport functions based on anatomic features (section 2.1.2, **Figure 3**). Thus, we could verify directly that all cross sections of fine roots corresponded to the morphology of absorptive roots according to McCormack et al. (2015). Furthermore, the distinction between intact roots and those with developed or initial symptoms of decay is easily performed on the cross sections. Using undisturbed soil samples and epifluorescence microscopy, we were able to quantify the



shows the *p*-value of the Dunn test for the different medians.

absorbing root surface and associated fungi. The method reliably differentiates between hyphae forming a mantel or EM. Based on our results it was possible to calculate the total absorbing root area root-fungi associations, which is a crucial component of plant-uptake models (Schnepf and Roose, 2006).

# 4.2. Controls of Rooting Density and Abundance of Root Classes

Our results confirm that beech trees adapt their root traits according to P availability, as shown by the efficient way of mycorrhization. This leads to a generally higher absorbing area at the P-poor site compared to the P-rich site, which compensates for the low P fluxes from soil to root surface. Soil texture, pHvalue, and other soil chemical parameters can have an influence on fine roots, so many studies address this aspect. In order to test these influences, a correlation of the fine root length density and the absorbing fine root surface per site with the soil chemical parameters (Ca, Fe, K, Mg, Mn, Na) as well as with the soil texture and pH-value was performed. We could only observe a positive correlation between root length density and soil Pcontent (**Figures S2, S3**). This indicates that P could have a significant influence on the root length density. However, we cannot claim that only P has an influence on the root length density. The significant negative correlation between the soil pH value and the fine root length density could be linked to an



**FIGURE 7** | Depth distribution of the root surface area index (RAI) (<2 mm)  $[m^2 m^{-3}]$  at BBR and LUE separately for the analyzed horizons. The fine root surface area down to 1 m soil depth related to ground area  $[m^2 m^{-3}]$  was estimated based on the thickness of horizons and is presented as cumulative bar. For LUE the Of horizon was added in fine root surface area index. For error bars see **Figure S4**. The mean root length density and standard deviation.

indirect correlation, an inter-correlation with the  $PO_4$  content. Thus, it can be excluded that the pH value causes the site specific differences between these two sites. This is in line with Richter et al. (2007, 2013) who show that there is an influence of base saturation and pH values on the fine roots of beech, but that this influence is smaller than expected. Now both LUE and BBR can be assigned to the base saturation group >5%, since for example the AhE horizon of LUE has a base saturation of 7% and the Ah horizon of BBR of 20%.

In our study  $RLD_{<1} mm$  was significantly higher at the P-poor site LUE than at the P-rich site BBR, whereas  $RLD_{1-2} mm$  showed no significant difference. The maximum ratio between  $RLD_{<1} mm$ and  $RLD_{1-2} mm$  was two to six times higher at the P-poor site than at the P-rich site.

Thus, the soil profile scale shows that the surface area and mycorrhization of fine roots (<1 mm) increase with P availability. The significant correlation between the ratio of mycorrhizal RLD to citrate-extractable  $PO_4$  at both sites (Figures S2, S3) indicated that root growth and mycorrhization is concentrated on horizons with high P availability. In contrast, comparing different sites, Fitter (1985) observed increasing specific root length with decreasing P availability. This is in agreement with our results regarding the comparison of the two different sites. At the P-poor site, LUE, the root length density is much higher than at the P-rich site BBR. Yet, the highest RLD<sub><1 mm</sub> was found at both sites in the two Oa horizons with the highest concentration of available P (Table 1, Figure 8). This statement is consistent with the observations of with Meier and Leuschner (2008), Meinen et al. (2009a,b) who showed that beech has a flat root system with the majority (approx. 75%) of fine roots in the Oa and in the topsoil (Meier and Leuschner, 2008; Meinen et al., 2009a,b) this is especially true for P-poor sites (Jonard et al., 2015). In agreement, in our study, the depth distribution of fine roots clearly differed at both sites: LUE had most of the fine roots in the forest floor and topsoil horizon, with 88% of the total fine roots occurring in the upper 11 cm of the soil (Figure 4), whereas



different mycorrhizal root lengths are given as numbers.

the root length density of the Cw horizon was much higher at BBR than at LUE (**Table 3**).

Further, the total mycorrhizal root length density is much higher in BBR the P-rich site than at the P-poor site LUE. According to Tatry et al. (2009) uptake rates in plant parts with ECM are two to three times higher than in root parts without mycorrhizal fungi, which supports the hypothesis that the greater proportion of mycorrhizal root length is found at the P-poor site. Also, Jones et al. (1990) showed that P inflow rates of ECM roots are about two times higher than those of non-mycorrhizal roots. Furthermore, mycorrhizal fungi can store P (Harley, 1989; MacFall and Slack, 1991) and thus possibly buffering phases of P starvation, which may also reflect an adaptation to low P availability. The low degree of mycorrhization in the Oa horizon of BBR and Cw of LUE seems to contradict the 100% mycorrhization for these sites reported by Zavišić et al. (2016). However, their results exclusively referred to the root tips, while our topological cross-sectional method examined the degree of mycorrhization over the entire root length and thus did not contradict his data. Mycorrhizal fungi can extend their RAI and their spatial coverage by forming EM. According to Nylund and Wallander (1992), this is promoted by low P contents. Therefore, the high frequency of mycorrhizal fungi and the high share of extramatrical mycelium, might help sustain sufficient P uptake at the site LUE in spite of low P availability.

The share of melanine and hyaline mycelia is different for different sites and horizons. Nosanchuk and Casadevall (2003) showed that melanin plays a protective role in unfavorable environments. One function of melanin could be the tolerance to water stress (Fernandez and Koide, 2013) and the protection against dehydration through storage of melanin in cell walls that in turn reduces water loss (Kogej et al., 2006). In contrast to hyaline cells, melanine cells have a high biosorptive capacity for numerous ions, which improves plant growth for plants in toxic environments (Fogarty and Tobin, 1996). Also, metal ions bound to melanin protect against toxicity and inhibition of extracellular hydrolytic enzymes (Fogarty and Tobin, 1996). This enables melanin to prevent enzymatic degradation (Nosanchuk and Casadevall, 2003) and makes hyphae more resistant to decay (Fernandez and Koide, 2013). Consequently, these properties increase the lifespan of melanine roots, thereby maintaining the P supply from litter or mineral soil as needed. The differences between both sites can be explained by the rough harsh soil chemical and physical soil properties at LUE with its higher acidity and sandy texture that could trigger water and mechanical stress. In BBR with its higher pH and water holding capacity, the environment for hyphae is probably more favorable. The higher robustness of melanine hyphae could also explain the strong dominance of the melanine hyphae in the EM. Extramatrical mycelium is more vulnerable to extreme conditions than hyphae in the mantle around the root, which is where we observed more hyaline hyphae. Finally, we cannot provide any information on mycorrhizal species composition, since our fine root classes are based on simple morphological characteristics. Such data have been provided for the study sites by Zavišić et al. (2018). Their molecular genetic determination of mycorrhizal fungi identified 14 different fungal morphotypes with 25 taxa by ITS sequencing. He showed that both sites had different taxa, with the greater biodiversity being found at the P-rich site indicating that different morphotypes might be linked to different species composition.

## 4.3. Root-Mycorrhizal Fungi Associations and P Nutrition Strategies of Beech Forest Ecosystems

As hypothesized we found clear differences between the traits of root/mycorrhizal associations at both sites, which would not have been identified by routine methods for fine root quantification: We found higher RLD on the P-poor than on the P-rich site and a higher portion of mycorrhizal root surface. Within the soil profiles, the plants clearly concentrate non-mycorrhizal roots in the P-rich horizons. In general, the different root morphology would support the different P nutrition strategies assumed for P-poor and P-rich beech forests (Lang et al., 2017). At the Prich site BBR where the forest ecosystem is assumed to acquire P from mineral resources we observed a high percentage of overall root surface in the Cw subsoil horizon besides the Oa horizon. This intense rooting in the subsoil seems to be mirrored by the concentration of soil organic matter at 1 m soil depth. The Corg content at BBR is still 11 mg C  $g^{-1}$  at this depth, while at the site LUE we only measured 1 mg C  $g^{-1}$ . This might indicate, that small fine roots neglected by conventional approaches represent a major source for deep-soil organic matter. In contrast, root length density was extremely high in the Oa and the AhE horizon in the P-poor ecosystem LUE, which is assumed to sustain its P supply by taking up P mobilized from organic matter. The prominent role of the Oa horizon for P uptake at the site LUE is further supported by its extension. Forest floor mass at LUE was five times higher than at site BBR.

## **5. CONCLUSION**

The results of our quantitative micro morphological cross sections analyzes of undisturbed soil samples, are in agreement with the idea that we could show that beech forest nutrition strategies are reflected in the distribution of fine roots and

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Ad-hoc-AG Boden (2005). Bodenkundliche Kartieranleitung: Ed.: Bundesanstalt für Geowissenschaften und Rohstoffe in Zusammenarbeit mit den Staatlichen Geologischen Diensten. E. Schweizerbart'sche Verlagsbuchhandlung (Nägele und Obermiller), 5th Edn. Hanover. the associated mycorrhization of European beech forests. The method gives new insights into the distribution and morphological traits of absorptive roots in forest stands including the size of the RAI and the degree and type of mycorrhization. Overall our study indicates, that the major part of fine roots had been overseen by conventional approaches for fine root quantification. Based on the detailed root information provided to confine idea of adaptation to forest ecosystem on P scarcity, which has to be validated by future studies: The depth distribution of the fine roots was controlled by P availability. Intense root growth in regions with high P availability, is compensated by reduced root growth in areas with low P availability. This mechanism would enhance the realization of either recycling or acquiring by European beech. As a consequence, the contrasting overall high rooting density at P-poor sites and the overall low rooting density at high-P sites is unlikely to be directly triggered by soil solution concentrations but seem to be rather the result of long-term adaptation to P scarcity.

## DATA AVAILABILITY STATEMENT

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

## **AUTHOR CONTRIBUTIONS**

CL carried out the experiment, analyses, and wrote the first draft of the manuscript. CL and HS-K performed the statistical analysis. All authors contributed to manuscript revision, read, and approved the submitted version.

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## SUPPLEMENTARY MATERIAL

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**Conflict of Interest:** The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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