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# Variation in root morphology amongst tree species influences soil hydraulic conductivity and macroporosity

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## Abstract

Natural approaches to flood risk management are gaining interest as sustainable flood mitigation options. Targeted tree planting has the potential to reduce local flood risk, however attention is generally focused on the hydrological impacts of catchment afforestation linked to generic tree features, whilst the species-specific impacts of trees on soil hydrology remain poorly understood. This study compared effects of different tree species on soil hydraulic properties. Monocultures of *Alnus glutinosa* (common alder), *Fraxinus excelsior* (European ash), *Fagus sylvatica* (European beech), *Betula pendula* (silver birch), *Castanea sativa* (sweet chestnut), *Quercus robur* (English oak) and *Acer pseudoplatanus* (sycamore maple) were used to determine effects of tree species identity on soil hydraulic properties (near-saturated K and soil water retention) in a sandy loam soil, North Wales, United Kingdom. The interaction of *F. excelsior* root properties and soil class on hydraulic conductivity was also examined in four different soils (Rendzic Leptosol, Haplic Luvisol, Dystric Fluvic Cambisol and Dystric Gleysol) across England and Wales. Fine root biomass (FRB) and morphological characteristics were determined at three depths (0-0.1, 0.1-0.2 and 0.2-0.3 m) and complemented by *in situ* surface measurement of soil hydraulic conductivity. Root morphological traits were closely associated with species identity and pore-size distribution, and FRB was strongly correlated with soil hydraulic conductivity ( $R^2=0.64$  for 0-0.1 m depth FRB;  $R^2=0.69$  for 0.1-0.2 m depth FRB). Fine root biomass of *F.*

*excelsior* was six-fold greater than *C. sativa* ( $p < 0.001$ ), and the frequency of 0.01 mm radius soil pores under *F. excelsior* was twice that of *Q. robur*. Near-saturated hydraulic conductivity under *F. excelsior* was  $7.91 \pm 1.23$  cm day<sup>-1</sup>, double the mean rate of the other species. Soil classification did not significantly influence FRB ( $p = 0.056$ ) or near-saturated hydraulic conductivity ( $p = 0.076$ ) in the 0.0-0.1 m depth soil, but soil water retention varied with depth. Species-specific traits of trees should be considered in landscape design to maximise the local hydrological benefits of trees.

**Keywords:** Land use; Infiltration;; *Hymenoscyphus fraxineus*; Hydrology; Soil porosity; Soil classification.

## 1. Introduction

Anthropogenic activities are driving an acceleration of climate change and, as a result, the occurrence and intensity of extreme weather events is predicted to increase (IPCC5 WGII, 2014). Precipitation in the United Kingdom (UK) over the past 250 years has increased during the winter and decreased during the summer (Dadson et al., 2017). Climate change has motivated greater attention to mitigating the impact of extreme events, such as flooding, with a policy focus on the role land use management can play (Mcintyre and Thorne, 2013).

Trees have the potential to influence soil hydrological processes by increasing water infiltration into soil, evapotranspiration, interception and groundwater recharge (Dadson et al., 2017; Dixon et al., 2016; Lane, 2017; Wolton et al., 2014). Plot-scale research has notably found that even when young (5-years-old), trees can increase infiltration rate by 67 times and reduce surface runoff by 78% compared with grazed pasture (Marshall et al., 2013), but heterogeneity of effects on hydraulic conductivity at plot scale is also evident (Chandler and Chappell, 2008). The interplay between soil and vegetation shapes soil hydraulic functions, but the relative importance of these functions is context specific. In arid zones, vegetation is highly influential in increasing hydraulic conductivity (Thompson et al., 2010), whereas soil class dominates the process in humid tropical and temperate (Geris et al., 2015) ecosystems. In contrast, soil classification has generally been shown to have little effect on

infiltration capacity, with interactions between soil fauna (e.g., earthworms), roots, plant species richness and soil structure of greater importance (Fischer et al., 2015; Jarvis et al., 2013).

The role of tree roots in shaping hydraulic response has often been overlooked (Chandler et al., 2018). Understanding of inter- and intra-species variation in root morphology is largely based on the questionable assumption that root architecture and hydrological function can be predicted from above-ground morphological characteristics (Sinacore et al., 2017). Therefore, a more thorough investigation of species-specific, below-ground hydrological function is required. Macropores within soil can be associated with root channels developed through the process of root production and turnover, and enable preferential flow (Ghestem et al., 2011). Bioturbation from soil flora and fauna can also increase porosity and hydraulic function, the effects of which are influenced by landuse intensity and antecedent soil conditions, such as pH (Spurgeon et al., 2013). Preferential flow in wooded ecosystems has been shown to be related to tree species; Luo et al. (2019) reported that coniferous forests dominated by *Platycladus orientalis* (L.) Franco (oriental arbo-vitae) exhibited greater preferential flow than deciduous forests dominated by *Quercus variabilis* Blume (Chinese cork oak). Separately, a positive relationship ( $R^2 = 0.91$ ) was identified between macroporosity and tree roots of *Pinus coulteri* D.Don (Coulter pine), but total porosity (and near-saturated conductivity) was greater under *Quercus dumosa* Nutt. (California scrub oak) and *Adenostoma fasciculatum* Hook. & Arn. (chamise), where conditions were more conducive to macrofaunal (e.g., earthworm) activity (Johnson-Maynard et al., 2002). Luo et al. (2019) reported that whilst tree roots were strongly associated with macropore development and preferential flow, the interaction between macroporosity, total porosity and infiltration was less clear. Soil total porosity and infiltration rate can have a positive relationship (Sun et al., 2018), however Bodner et al. (2014) attributed an increase in infiltration to an increase in macroporosity in soil where total porosity remained unchanged. Inconsistent effects describing the relationships between total porosity, macroporosity, preferential flow and tree species identity implies that more research is required to understand these associations.

77 Preferential flow in the vadose zone mediates water infiltration and is associated with macropores,  
78 including artificial drainage (Bathurst et al., 2018; Marshall et al., 2009), macrofaunal pathways  
79 (Bargues Tobella et al., 2014), biomat flow (Gerke et al., 2015) and root channels (Zhang et al., 2015).  
80 However, not all fine roots are conduits for preferential flow (Luo et al., 2019), suggesting that root  
81 size distribution may be more important than root biomass. For example, root length density has been  
82 shown to have a strong positive correlation with preferential flow (Zhang et al., 2015) but this  
83 relationship is spatially variable (Luo et al., 2019).

84 The difference in fine root production across a spectrum of the broadleaved tree species that are  
85 abundant in Europe, and the consequential effect on soil hydraulic conductivity, is largely unknown.  
86 Fine root production is known to be plastic, with its spatial distribution being highly responsive to  
87 antecedent moisture conditions (Fan et al., 2017), which is influenced by soil texture as well as by  
88 climate. Differences in hydrological response have been shown between coniferous and deciduous  
89 forest ecosystems, but the response was mitigated by spatially contrasting soil texture (Luo et al.,  
90 2019). The relative influence of tree species identity and soil classification on infiltration capacity  
91 remains poorly understood. The aim of this study was to characterise the root morphology of seven  
92 species of broadleaved, deciduous trees and investigate the relationship with near-saturated soil  
93 hydraulic conductivity within one type of soil. The study then seeks to understand the effect of soil  
94 classification on hydraulic conductivity in plantations of a single using tree species. The objectives  
95 were to (i) investigate the variation in infiltration rate between seven tree species growing in the same  
96 soil classification and (ii) compare the tree species' corresponding root morphological characteristics  
97 to determine whether soil hydraulic function depends on species' root characteristics, then (iii) to  
98 investigate the relative influence of tree roots and soil classification on soil hydraulic function. We  
99 hypothesise that (i) tree species affect soil hydraulic conductivity; (ii) tree species' growing on the  
100 same soil differ in their production of fine root biomass (FRB) and infiltration rate; and (iii) soil  
101 classification affects the soil hydraulic function associated with the abundant European tree species  
102 *Fraxinus excelsior* L (European ash).

## 2. Methods

### 2.1 Site descriptions and experimental design

The BangorDiverse forest diversity and ecosystem function experiment, located at the Henfaes Research Centre, Abegwynnregyn, UK (53°14'15"N, 4°1'4"W), was used to determine the effect of tree species on soil hydraulic function. Monocultures of seven tree species were planted as 1.0 m tall saplings in March 2004: *Alnus glutinosa* [L.] Gaertner (common alder), *F. excelsior*, *Fagus sylvatica* L. (European beech), *Betula pendula* Roth. (silver birch), *Castanea sativa* Mill (sweet chestnut), *Quercus robur* L (English oak) and *Acer pseudoplatanus* L. (sycamore maple) (Ahmed et al., 2016). Initial planting density was 10,000 stems ha<sup>-1</sup>, but trees were thinned to 2,500 stems ha<sup>-1</sup> in 2012/2013 to facilitate continued tree development. Randomised, replicate plots (0.1 ha) of each species (n=4) were blocked across two adjacent fields (2.36 ha total area). The soil at BangorDiverse is a Dystric Fluvisol Cambisol, developed from glaciofluvial deposits (Smith et al., 2013) with pH ranging from 5.4 (surface) to 6.3 (1-m depth) (Ahmed et al., 2016). Soil texture is a sandy loam/loam determined by laser diffraction (Coulter LS particle size analyser) from soil in the 0-0.1 m depth. The site is hyperoceanic with mean annual rainfall of approximately 950 mm and mean annual air temperature of 10.6 °C (Gunina et al., 2017).

Plots of *F. excelsior* planted in different sites across the UK with four contrasting soil classifications (IUSS Working Group WRB, 2015) were used to investigate how interaction of a single tree species with soil classification influences soil hydraulic function. Originally established as part of a provenance trial in 1993 (Cundall et al., 2003), three sites, Gloucestershire (England), Hampshire (England) and Gwynedd (Wales), were selected based on the soil classifications (Table 1) that best represented the range of textural characteristics (sand, silt or clay) commonly occurring across the UK. Each experimental site consisted of three fully replicated, randomised blocks of different provenances of *F. excelsior*. Saplings (same age from seed) were planted at 2500 stems ha<sup>-1</sup> and had subsequently been thinned to 50% density at the Gloucestershire site only. One plot from each block (n=3) comprised of

*F. excelsior* trees of UK (Powys, Shropshire) or French (Normandy) provenance were selected for study. Due to the presence of the fungal pathogen *Hymenoscyphus fraxineus* at the Hampshire site only, plots were selected where only visibly healthy trees were found following condition assessment (SI 1). To increase the diversity of soils used in this analysis (Table 1), an additional site with *F. excelsior* (provenance unknown) established in 1987 at Rothamsted Research, North Wyke, Devon (England) was selected. At North Wyke, three plots were randomly selected from two blocks, avoiding edge trees. All plots were planted with seedlings at 2500 stems ha<sup>-1</sup> and had not been thinned. No obvious signs of *H. fraxineus* were present at North Wyke.

## 2.2 Root morphology

Two soil cores of 0.08-m diameter were collected from three depths (0-0.1, 0.1-0.2, 0.2-0.3 m) equidistant between two trees randomly selected near to the centre of each plot to mitigate against edge effects (SI 2). Roots were collected to a depth of 0.3 m; in soil above this depth, in temperate forest ecosystems, 65% of roots exist and there is a predominance of fine ephemeral roots involved in nutrient and water uptake (Jackson et al. 1996). To minimise canopy damage and variation introduced by root growth during the sampling period, 168 samples were collected between January and February, after leaf fall and during a period of dormancy in line with previous sampling campaigns conducted at the site (Smith et al., 2013). Soil cores were placed into sealable polythene bags and stored at 4 °C for a maximum of 4 days before processing.

Each core was washed with water in a sieve stack (1- and 2-mm mesh size) to remove soil adhered to roots and separate roots into two size classes, fine (<2 mm diameter (ø)) and coarse (>2 mm ø), the latter of which were discarded. Tree species identity of the roots was based on morphological characteristics, such as surface colour, structure and colour of the periderm and ramification pattern, outlined by Mrak and Gricar (2016) and necromass (dead fine roots) was identified based on black or dark brown colour and a decaying fragmented appearance (Eissenstat and Yanai, 1997; Leuschner et

al., 2004 ; Smith et al., 2013). Fine roots were scanned using an Epson 4990 scanner at a resolution of 300 dots per inch (dpi) and images were analysed with WinRhizo (version 2005c, Regent Instruments Inc., Quebec, Canada) to measure fine root length, surface area, surface volume, projected surface area and number of root tips, divided into 20 (0.1 mm) diameter classes (0-2 mm). Necromass and the biomass of fine and coarse roots were determined after drying at 80 °C until constant mass. Data from the two soil cores collected per plot were averaged to avoid within-plot pseudoreplication.

### *2.3 Root characteristics*

Root area index (RAI,  $\text{m}^2 \text{m}^{-2}$ ) was derived from the root surface area divided by the surface area of the sampled core. Specific root area (SRA,  $\text{m}^2 \text{kg}^{-1}$ ) was calculated from the surface area of fine root divided by root dry mass (Lohmus et al., 1989). Specific root length (SRL,  $\text{m g}^{-1}$ ) was determined from the total length of fine roots divided by root dry mass (Ostonen et al., 2007). Root length density (RLD,  $\text{cm cm}^{-3}$ ), which indicates the proportion of soil occupied by fine roots, was estimated from the ratio of root length to the volume of the sampled core. Root tip density (RTD) was calculated as thousands of tips per  $\text{m}^2$ . For each of the aforementioned root metrics an arithmetic mean was calculated from data exported from WinRhizo output.

### *2.4 Soil hydraulic function*

Minidisk infiltrometers (0.045 m  $\varnothing$ ) (Meter Group, Pullman, USA) were used to measure the rate of infiltration of water into soil and to calculate near-(field)-saturated hydraulic conductivity ( $K_{fs}$ ) within each plot. Surface vegetation was carefully removed, and a thin layer of fine sand ( $\sim 0.001$  m) was applied to the soil surface to ensure optimal contact between the infiltrometer disc and the soil. The tension was set at -0.02 m to eliminate water flow through the largest macropores ( $> 0.742$  mm), to provide a more representative estimation of water flow through the soil matrix and to achieve steady-state infiltration rate. Water level was recorded every minute until 20  $\text{cm}^3$  of water had infiltrated the soil. Three measurements were taken at each plot to give an average  $K_{fs}$ . Near-saturated hydraulic



conductivity for the respective soil water potential was calculated using the method of Zhang (1997) and van Genuchten soil classification tables (Meter Group Inc, 2018).

At each plot, a 250 cm<sup>3</sup> soil core was collected from the 0-0.05 m and 0.1-0.15 m depths. Cores were stored at 4 °C and then soaked for at least 24 hours in degassed, deionised water prior to analysis. Soil water retention was measured using a HYPROP 2 (Meter group, Pullman, USA) (Schindler et al., 2010), and then dry bulk density and porosity were determined for the cores (SI 3). To account for the stoniness of the experimental plots, stones (>0.002 m  $\phi$ ) were sieved out of the oven-dried soil and weighed (SI 3). The vapour equilibration technique (Scanlon et al., 2002) was used to measure the dry-end matric potential on sub-samples taken from each core. Soil water retention curves (SWRC) were modelled using the HypropFit (Schindler et al., 2010) (UMS, Munich, Germany) implementation of the Fredlund-Xing water retention model (Fredlund and Xing, 1994), using the measured soil water retention, dry bulk density,  $K_{fs}$  (applied to cores from 0-0.05 m depth only) porosity, dry-end matric potential, volumetric moisture content and stoniness data.

Effective soil pore-size distribution was estimated using the method outlined by Blonquist et al. (2006). Hydraulic capacity was estimated using data from the SWRC (modelled in HypropFit) to derive the change in moisture over the change in hydraulic head ( $d\theta_v/dh$ ). Hydraulic capacity was plotted as a function of pore radius. The scaled effective pore-size distribution associated with each tree species was then derived by taking the inverse relationship between pressure (h) from the water retention curve and log10 pore radius, resulting in a dimensionless, scaled, effective pore-size distribution. The distribution is displayed as a function of effective pore radius  $f(r)$  proportional to the abundance of each pore-size within a given volume of soil.

To give context, *in situ* soil moisture was measured using ML3 ThetaProbe Soil Moisture Sensors (Delta-T Devices Ltd, Cambridge, UK) (n=9) in each plot at 10 cm depth. Particle-size distribution was ascertained using an air-dried sub-sample from soil used for the HYPROP analysis, repeatedly

quartered to mitigate selection bias (Lebron and Robinson, 2003) (Table 1). Particle-size distribution was determined from a 0.5-0.8 g subsample of sieved (<2 mm) soil using a LS13 320 laser diffraction particle-size analyser (Beckman Coulter Inc, Indianapolis, USA) (Table 1). Soil organic matter concentration was determined by loss-on-ignition (LOI) analysis of 10 g of sieved (<2 mm) soil (Ball, 1964) (Table 1). For quality assurance, two standard soil and two replicate samples were included for all LOI and particle size analyses.

## 2.5 Statistical analyses

Two statistical models were used to analyse the datasets: (i) for the data collected from BangorDiverse (n=4), a two-factor ANOVA to test factors and interaction effects, with species and depth as factors, and root biomass, root morphological characteristics and  $K_{fs}$  as dependant variables; (ii) for data collected at the pan-UK *F. excelsior* provenance trial sites (n=4), a two-factor ANOVA with soil classification and soil depth as factors and root biomass, root morphological characteristics and  $K_{fs}$  as dependant variables. The Tukey Honest Significant Difference (HSD) post-hoc test was used to determine within-factor significance for both statistical models. Relationships between dependent variables were explored using ordinary linear regression. All data were tested for homogeneity of variance using Levene's test and normality using the Shapiro-Wilk's test. Root biomass, root morphological variables and  $K_{fs}$  data were log transformed to satisfy normality. To visualise the relationships between variables, the dimensionality of the dataset was reduced from 44 parameters that included root morphological metrics (e.g., SRL, RAI, SRA, RLD) within three soil layers (0-0.1 m, 0.1-0.2 m and 0.2-0.3 m),  $K_{fs}$  at the soil surface and soil porosity within two soil layers (0-0.05 m and 0.10-0.15 m) by conducting a principal component analysis (PCA). Stepwise multiple regression (forward and backwards) was then used to determine the parameter that best predicted  $K_{fs}$ . All statistical analyses were completed with SPSS v22.0 (IBM SPSS, Armonk, NY, USA) with  $p < 0.05$  used as the limit for statistical significance. All figures were produced using SigmaPlot v13.0 (Systat Software, San Jose, CA, USA).

### 3. Results

#### 3.1 Tree species' effects

##### 3.1.1 Hydraulic conductivity and root biomass

Mean surface  $K_{fs}$  ranged from  $3.47 \pm 0.56$  standard error  $\text{cm day}^{-1}$  for *A. pseudoplatanus* to  $7.91 \pm 1.23$   $\text{cm day}^{-1}$  for *F. excelsior*, although the difference between the species did not reach the threshold of statistical significance ( $p=0.056$ ) (Fig. 1a). However, a positive correlation ( $R_{253,254} = 0.64$  (0-0.1 m depth) and  $R^2 = 0.69$  (0.1-0.2 m depth)) was observed between tree species' FRB and  $K_{fs}$ , with a high degree of variation around mean  $K_{fs}$  within some species (e.g. *Q. robur*).

Fine root biomass was affected by both species and soil depth ( $p<0.001$ ), but no interaction effect was evident (Table 2). *Fraxinus excelsior* was the species producing highest FRB at every soil depth (Table 3), with the largest difference in FRB between *F. excelsior* and the other species at a depth of 0-0.1 m ( $p<0.001$ ; Fig 1b). Fine root biomass of *F. excelsior* was between three-fold (*B. pendula*;  $p<0.001$ ) and six-fold (*C. sativa*;  $p<0.001$ ) greater than the other species. Deeper in the soil where the proportion of total *F. excelsior* FRB was much less (24%, 0.1-0.2 m; 17%, 0.2-0.3 m), *F. excelsior* FRB was greater than *F. sylvatica* FRB only ( $p=0.05$ , 0.1-0.2 m;  $p<0.01$ , 0.2-0.3 m). The biomass:necromass (B:N) ratio of *A. pseudoplatanus* (37.19) was significantly ( $p<0.05$ ) greater than *A. glutinosa* (4.21) and *C. sativa* (5.56) within the 0-0.1 m soil layer and was significantly greater for *A. pseudoplatanus* (103.9) than *A. glutinosa* (3.27), *C. sativa* (1.05), *F. sylvatica* (6.60) and *Q. robur* (5.23) within the 0.1-0.2 m soil layer ( $p<0.05$ ).

Soil total porosity (0-0.05 m depth; Fig 1c) was similar between all species ( $p>0.05$ ) at all soil depths. Despite the aforementioned positive correlation between  $K_{fs}$  and FRB, a similar relationship was not observed between FRB and total porosity. There is some evidence of a positive linear relationship between the mean  $K_{fs}$  and FRB of each species, although it reached the  $p<0.05$  threshold of significance in the 0.1-0.2 m depth only. Fine root biomass explained 64, 69 and 25% of the variation in  $K_{fs}$  for the 0-0.1 m, 0.1-0.2 m and 0.2-0.3 m depths, respectively (Fig. 2a-c).

### 3.1.2 Soil water retention and pore-size distribution

Saturated soil water content was highest for *F. excelsior* (57%) and lowest for *F. sylvatica* (52%) in the 0-0.05 m soil layer (Fig. 3a). As soil water potential decreased, the soil water content under *F. excelsior* decreased rapidly, becoming comparable to the other species. Continued decreases in soil water potential caused *F. excelsior* to have the second lowest retention capacity. Conversely, *Q. robur* was ranked 5<sup>th</sup> in species' retention capacity at saturation but retained the highest percentage of soil water content at mid-range potentials (i.e., between -100 and -1000 cm). *Castanea sativa* had consistently low soil water content compared with the other species. Within the 0.10-0.15 m soil layer (Fig. 3b), *Q. robur* had the highest water content (57%) at saturation, whereas *F. excelsior* had the second lowest (50%), with *F. sylvatica* lowest (49%). All species had similar water content once pressure was applied (< -10 cm), apart from *C. sativa*, which again had consistently lower soil water content than other species.

Figures 3c and 3d show the scaled effective pore-size distribution. Soil developed under *F. excelsior* exhibited the greatest abundance (0.24) of macropores, followed by *B. pendula* (0.20), whilst the pore-size distributions of soil under *Q. robur* and *C. sylvatica* are skewed towards smaller pore sizes. By contrast, the proportion of macropores deeper in the soil (0.1-0.15 m) were similar amongst species, with the exception of *Q. robur* and *A. pseudoplatanus* (Fig 3d).

### 3.1.3 Root morphological traits

Tree species and separately soil depth affected all root traits (both  $p < 0.001$ ) except soil depth for SRA ( $p > 0.05$ ), but there were no species  $\times$  depth interactions (Table 4). In the 0-1.0 m soil layer, *F. excelsior* had greater RLD ( $6.56 \pm 0.65 \text{ cm cm}^{-3}$ ,  $p < 0.05$ ) and RAI ( $6.02 \pm 0.86 \text{ m}^2 \text{ m}^{-2}$ ,  $p < 0.01$ ) than all other species, more than 11 times greater than lowest ranked *C. sativa* (RLD  $0.57 \pm 0.07 \text{ cm cm}^{-3}$ , RAI  $0.51 \pm 0.51 \text{ m}^2 \text{ m}^{-2}$ ). Root tip density (RTD) was also greatest in *F. excelsior* ( $1275.01 \pm 199.3 \times 10^2 \text{ m}^{-2}$ ), significantly more than *A. glutinosa* ( $242.89 \pm 45.28 \times 10^2 \text{ m}^{-2}$ ;  $p < 0.001$ ), *C. sativa* ( $174.35 \pm 17.17 \times 10^2 \text{ m}^{-2}$ ;  $p < 0.001$ ) and *Q. robur* ( $p < 0.01$ ) (Table 5). The lowest RTD, associated with *C. sativa*, was more than seven-fold less than *F. excelsior*. *Fraxinus excelsior* had the greatest RLD, RAI and RTD in the 0.1-

0.2 m soil layer ( $p<0.01$ ) and in the 0.2-0.3 m soil layer; RLD and RAI of *Fraxinus excelsior* ( $1.55 \pm 0.47$  cm cm<sup>-3</sup> RLD;  $1.78 \pm 0.57$  m<sup>2</sup> m<sup>-2</sup> RAI) were three- to eight-fold and four- to seven-fold greater respectively than all other species ( $p<0.05$ ) except *A. pseudoplatanus* ( $p>0.05$ ; Table 5).

Ordination analysis was used to examine the relationship between tree root morphological traits and soil physical properties developed under the different tree species. The dimensionality of the data was reduced to three principal components (PC) that explained 95% of the variation. Principal component 1 explained 63%, PC2 18% and PC3 14% of the variation (Fig. 4). Tree species were tightly grouped together along the dominant PC1 with the exception of *F. excelsior*, which was strongly separated and associated most strongly with FRB, RAI, RTD, RLD and root projected surface area. Necromass was weakly separated from other root traits along PC2 and associated with *A. glutinosa* and *F. excelsior* (Fig 4a). Soil porosity in the 0-0.05 m depth and  $K_{fs}$  were associated with each other along PC1 and weakly associated with *F. excelsior* and *A. glutinosa* along PC2 compared with the other five species, whereas, deeper in the soil (0.1-0.15 m), total porosity related more strongly to the other five species than *F. excelsior* and *A. glutinosa*. Fine root biomass and other morphological traits (i.e., root projected surface area, RAI, RTD and RLD) were all closely associated with each other along PC1, and with *F. excelsior*. Stepwise multiple regression analysis (forward and backward) showed that root necromass was the best single predictor of  $K_{fs}$  ( $R^2 = 0.224$ ;  $p<0.05$ ) with all other variables excluded during the analysis.

### 3.2 *Fraxinus excelsior* across soil classifications

Fine root biomass of *F. excelsior* differed significantly amongst soil depths and only between soil class where the fungal pathogen *Hymenoscyphus fraxineus*, which causes ash dieback disease on *F. excelsior* affected tree growth (i.e., Rendzic Leptosol). No interaction effect was evident between soil class and depth (Table 6). Fine root biomass was lowest in the Rendzic Leptosol through the whole profile (0-0.3 m), with a B:N ratio of 1.16, compared with the Haplic Luvisol (3.17), Dystric Fluvisol Cambisol (6.62) and Dystric Gleysol (2.04) soils (SI 4). The relationship between FRB and hydraulic conductivity

previously observed across all tree species was reproduced when the relationship between *F. excelsior* FRB and hydraulic conductivity was examined across the four soils; the  $R^2$  was 0.49 for the two soil layers 0-0.1 and 0.1-0.2 m and 0.43 for the 0.2-0.3 m layer.

Soil water retention curves were similar under *F. excelsior* across all four soil classifications in the surface layer (Fig 5a). At saturation, the soil water content at 0.0-0.05 m depth did not vary significantly and ranged between 61% and 57% for all soil classes. The shapes of the retention curves were also similar throughout the range of water potentials. Conversely, SWRC from deeper in the soil profile (0.1-0.15 m) differed substantially (Fig 5b). While the SWRC of the Dystric Gleysol from Devon retained the same form as the surface soils, all other soils had decreased water retention at saturation with depth. The two silty clay loam soils, Haplic Luvisol and Rendzic Leptosol, had the greatest change in soil water content at saturation, both reducing from ~58% at the surface to ~48 and ~42% respectively with depth. The silt loam, Haplic Luvisol, soil had a unimodal pore-size distribution, but the other soils all had a bimodal distribution (Fig 5c&d). For all four soils macro- and meso-size pores were clearly evident in the surface layer, but decreased with depth, particularly the mesopores, with small pores becoming more prevalent with depth especially in the Dystric Gleysol and Haplic Luvisol soils.

Compared with reference soils in the Rosetta database (Schaap et al., 2001), soils from the present study retained a greater volume of water at saturation, regardless of soil classification (Fig. 6). The modelled soil water retention, based on physical soil characteristics of agricultural soils, was 15-50% less at saturation than those measured in the forested soils of the present study. Increasing soil water potential (-cm) rapidly reduced the volumetric water content of measured SWRCs to become comparable with the predicted reference soils by -100 cm.

## 4. Discussion

### 4.1 Tree root morphology and hydrology

This study showed that FRB production is tree species-specific, broadly agreeing with Chandler et al. (2018). Notably, *F. excelsior*, a ubiquitous species across much of Europe, establishes fine roots far more extensively, up to six-fold greater biomass, than the other common European broadleaved species assessed. Across species, total soil porosity remained consistent, but variation in FRB changed soil macroporosity and soil water retention. The results indicated that, although variation in species' FRB roughly mirrors that of  $K_{fs}$ , there was no relationship between FRB and total soil porosity. Soil under *F. excelsior* had the greatest water retention capacity at saturation (soil water potential = -1 cm), but the negligible variation between species indicated comparable total porosity. As soil water potential decreased the soil water content generated from soil collected under *F. excelsior* decreased rapidly, signifying the low bulk density and larger pore sizes (Radcliffe and Simunek, 2010) associated with *F. excelsior*.

Differences in pore-size distribution, rather than total porosity, linked to tree species-specific differences in fine root morphology are likely to be driving the relationship between tree species and hydraulic conductivity, but is moderated by fine root necromass. *Fraxinus excelsior* had the largest  $K_{fs}$ , root biomass and number of macropores, but the overall total porosity did not differ significantly from the soils under the other six tree species. The high FRB of *F. excelsior* might suggest adventitious root development and a greater RTD leading to the creation of macropores surrounding the root (Ghestem et al., 2011). However, it is apparent that FRB, projected root surface area and RTD are not as strongly related to porosity as are other root traits (Fig. 4). Despite the high FRB and hydraulic conductivity associated with *F. excelsior*, a correspondingly high RTD was not identified, suggesting that RLD, rather than RTD, is an important factor in the creation of macropore channels.

Despite nuanced relationships between live root morphological variables, macroporosity and  $K_{fs}$ , root necromass was the best predictor of  $K_{fs}$  suggesting that root turnover has an important role in soil hydraulic function. Fine root longevity in trees is complex, ranging from days to years (Bengough, 2012) and is dependent on root diameter, root density, nitrogen concentration, colonisation of

mycorrhizal fungi and phenolic compound accumulation mediated by interaction with soil fauna (Eissenstat et al., 2000). During root development, exuded organic compounds contribute to the stability of the root channel, but following root death dehydration initially occurs, allowing gradual decomposition that creates progressively larger channels within the soil matrix available for preferential flow, and subsequently sub-surface sediment transfer causes channels to collapse or fill over time (Bengough, 2012; Ghestem et al., 2011). Variation in root turnover rates should have a large influence on the size and longevity of root-derived macropores (Wang et al., 2020).

Tree root morphological traits in this study better explained  $K_{fs}$  variations near the soil surface (0-0.2 m) than deeper in the soil (0.2-0.3 m depth). Root length density was greatest near the soil surface facilitating connectivity of root-induced macropores and greater opportunity for infiltration. A similar strong relationship between macroporosity near the soil surface and preferential flow in three tree species (*Styphnolobium japonicum* (L.) Schott, *Platycladus orientalis* (L.) Franco, and *Quercus dentata* Thunb.), which diminished with depth, has also been reported (Zhang et al., 2015). A comparison with pedotransfer functions, largely used for agricultural soils, indicated that, by excluding sub-surface flow through macropores, hydraulic functions quickly converge with those predicted by the pedotransfer functions for the given soil texture in the 0-0.1 m soil layer. However, deeper in the profile (0.1-0.3 m) where the density of fine roots is lower, soil texture had a greater influence on soil hydraulic conductivity. Results of the present study, combined with the apparent lack of accountability for macropores generated by trees in pedotransfer functions, suggests that improvement could be made to the parameterisation of hydrological models based on the below-ground characteristics of vegetation.

#### 4.2 Soil classification and hydrology

Our study explored whether variation due to soil textural properties would temper the influence of afforestation with a single tree species (*F. excelsior*) on water retention capacity. Data from the forested plots were compared to agricultural soils with the same textural properties (loam, silty clay



loam and clay loam) in the Rosetta database to obtain comparable values of hydraulic response. Modification of soil structure by the presence of trees enabled greater water retention capacity at saturation (Fig. 6). As water potential increased, which effectively excludes the influence of macropores, the forest SWRCs migrate closer to the Rosetta predictions. Therefore, landcover, specifically the presence of trees, appears to mediate the influence of soil textural properties on hydraulic response, regardless of soil classification, although within landcovers impacts, such as tillage, may regulate the response.

Using *F. excelsior* as an example, this study showed that soil texture, a considerable influence on ambient soil moisture, does not influence fine root growth near the soil surface. Furthermore, in the 0-0.1 m depth, fine tree roots modified pore-size distribution, negating the effect of soil class on hydrological function. At 0-0.1 m depth, where 50-58% of total FRB of *F. excelsior* was present, little variation in soil water retention was observed between sites differing in soil classification. Deeper in the soil, soil water retention was more divergent amongst sites as the influence of fine roots decreased and soil class started to dominate the hydraulic response. Hydraulic conductivity, therefore, is influenced by the combination of root morphology and soil classification, which varied with depth. Indeed, within-species variation in root morphology and rooting extent throughout the soil profile has been shown to be contingent on ambient hydrological soil conditions, oxygen availability and access to groundwater resources (Feng et al., 2017).

During very dry conditions, such as those recently preceding the study period (mean volumetric soil water content of 16%), soil class had a nuanced effect on rooting morphology and macroporosity. The sandy silt loam and clay loam textures of the Dystric Fluvic Cambisol and Dystric Gleysol exhibited similar pore-size distributions. By contrast, the silty clay texture of the Haplic Luvisol was associated with a lower FRB. There was a lack of organic matter, or limestone, in the Haplic Luvisol that could disaggregate the clay compared with the other clay-containing soils (i.e., Dystric Gleysol and Rendzic Leptosol). The high clay content resulted in a substantially hardened soil that reduced plasticity and

was likely to be related to the observed lower abundance of macropores. Root dieback, however, caused by tree disease may have a greater, though time limited, impact on soil hydraulic function than soil classification. Root dieback is positively associated with crown reduction due to infection from *Hymenoscyphus fraxineus* (Bakys et al., 2011). Where *H. fraxineus* was observed at a moderate - advanced stage (assessment methods described in SI) (i.e., Hampshire; Rendzic Leptosol), necromass accounted for half of the total fine root mass (B:N = 0.98) in the 0-0.1 m depth, substantially more than in the other clay-dominant soils (Haplic Luvisol, 2.49; Dystric Gleysol, 2.05). Once necromass has fully decomposed, the residual root channels will be vulnerable to collapse, potentially reducing hydraulic conductivity in the longer-term.

#### 4.3 Implications for land managers

*Fraxinus excelsior* had the greatest potential to improve surface water infiltration regardless of soil class. A ubiquitous species in much of Europe, *F. excelsior* is likely to have a disproportionately larger influence on landscape hydraulic function than the other tree species assessed here due to its root morphology and influence on macroporosity.. Therefore, loss of *F. excelsior* in the landscape due to the fungal pathogen *H. fraxineus* could have serious implications for local soil hydrological function throughout Europe. Consideration of hydraulic function should be a major component in the selection of alternative tree species to replace *F. excelsior*, and tree species' root morphological traits and influence on soil hydrology should be used as a criterion to select tree species in the future to maximise the potential benefits of establishing new woodlands. However, whilst results of the present study showed that tree species-specific root morphological traits have a role in altering soil hydraulic function at the plot scale, the complex interactions that influence catchment hydrology (e.g., field boundaries, land use and drainage) suggest that caution should be exercised before extrapolating such plot-scale results to the landscape scale.

## 5. Conclusion

Species-specific variation in fine root morphological characteristics of seven common European broadleaved tree species were shown to alter soil macroporosity and hydraulic function. Fine root length density and necromass were correlated with an increased abundance of macropores within the soil, facilitating greater hydraulic conductivity, despite little change in total porosity. Notably, *F. excelsior* had up to a six-fold greater FRB than the other tree species studied, however RLD rather than FRB was shown to be the strongest driver of the observed changes in macroporosity.

Soil water retention curves and porosity data indicated that tree roots influence soil structural characteristics in the 0-0.1 m layer of the soil, where more than 50% of the FRB was present, maximising macroporosity regardless of soil texture. Species with the greatest RLD exhibited correspondingly greater macropore abundance and higher hydraulic conductivity when soils were at or close to saturation.

The species-specific influence of trees on hydraulic function and the associated impact of tree diseases, such as the fungal pathogen *Hymenoscyphus fraxineus* which causes ash dieback disease on *F. excelsior*, suggests that changes to the composition of tree species present in the landscape could have implications for hydrological hydraulic regulation. Further work is necessary to determine if hydrological models can be improved by the incorporation of below-ground tree trait data.

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## 615 **Tables**

616 **Table 1.** Location of *Fraxinus excelsior* provenance trial experimental plots by World Reference Base (WRB) soil classification (WRB for soil resources, 2006),  
 617 UK county, mean particle size distribution of the mineral soil to 0.015 m soil depth and soil texture based on measured soil particle size distribution (Soil  
 618 Classification for England and Wales) (Avery, 1980).

WRB classification soil group	Site (UK county)	Latitude Longitude	Mean soil particle size distribution (%)			Soil texture	Mean soil organic matter content (%)
			Sand	Silt	Clay		
Dystric Fluvisol Cambisol	Gwynedd	53° 14' 19.38" N 4° 01' 05.91" W	40	44	16	Sandy silt loam	6.4
Haplic Luvisol	Gloucestershire	51° 54' 24.93" N 2° 18' 39.68" W	20	59	21	Silty clay loam	6.5
Rendzic Leptosol	Hampshire	51° 12' 02.02" N 1° 31' 39.48" W	16	57	26	Silty clay loam – limestone rich	7.0
Dystric Gleysol	Devon	50° 46' 12.14" N 3° 54' 08.79" W	25	51	23	Clay loam	11.5

619

620

621 Table 2. Main effects of seven tree species' (*Alnus glutinosa*, *Fraxinus excelsior*, *Fagus sylvatica*, *Betula pendula*, *Castanea sativa*, *Quercus robur* and *Acer*  
 622 *pseudoplatanus*) fine root biomass in three soil depths (0-0.1, 0.1-0.2, 0.2-0.3 m).

Factor	df	F	p-value
Depth	2	16.156	<0.001
Species	6	11.677	<0.001
Depth*Species	12	0.831	0.618

623

624 Table 3. Mean fine root biomass and rank order of seven tree species (*Alnus glutinosa*, *Fraxinus*  
625 *excelsior*, *Fagus sylvatica*, *Betula pendula*, *Castanea sativa*, *Quercus robur* and *Acer pseudoplatanus*)  
626 at 0.1 m depth intervals and for the whole sampled profile (0-0.3 m). Mean fine root biomass at each  
627 depth is given as a proportion of the whole profile (0-0.3 m) fine root biomass (%), B:N describes the  
628 biomass:necromass ratio. SE =  $\pm 1$  standard error. Superscript letters denote Tukey post hoc  
629 comparison ( $p < 0.05$ ) between species within each soil depth.

	Fine root biomass (kg m <sup>-2</sup> )		Rank order	Proporti on of total fine root biomass (%)	Necromass (kg m <sup>-2</sup> )		B:N ratio	
<b>0–0.1 m</b>	Mean	SE			Mean	SE	Mean	SE
<i>Alnus glutinosa</i>	0.84 <sup>b</sup>	$\pm 0.15$	4	60.31	0.25	$\pm 0.05$	4.21 <sup>b</sup>	$\pm 1.53$
<i>Fraxinus excelsior</i>	3.05 <sup>a</sup>	$\pm 0.40$	1	59.01	0.36	$\pm 0.14$	12.17 <sup>ab</sup>	$\pm 3.69$
<i>Fagus sylvatica</i>	0.55 <sup>b</sup>	$\pm 0.12$	6	51.36	0.03	$\pm 0.01$	16.85 <sup>ab</sup>	$\pm 3.69$
<i>Betula pendula</i>	1.12 <sup>b</sup>	$\pm 0.13$	2	48.71	0.14	$\pm 0.05$	10.08 <sup>ab</sup>	$\pm 2.13$
<i>Castanea sativa</i>	0.45 <sup>b</sup>	$\pm 0.09$	7	36.91	0.07	$\pm 0.04$	5.56 <sup>b</sup>	$\pm 2.81$
<i>Quercus robur</i>	0.62 <sup>b</sup>	$\pm 0.29$	5	49.10	0.12	$\pm 0.07$	7.30 <sup>ab</sup>	$\pm 2.51$
<i>Acer</i> <i>pseudoplatanus</i>	0.87 <sup>b</sup>	$\pm 0.28$	3	42.88	0.08	$\pm 0.06$	37.19 <sup>a</sup>	$\pm 16.73$
<b>0.1–0.2 m</b>								
<i>Alnus glutinosa</i>	0.26	$\pm 0.07$	7	19.04	0.15	$\pm 0.05$	3.27 <sup>b</sup>	$\pm 1.92$
<i>Fraxinus excelsior</i>	1.22	$\pm 0.31$	1	23.56	0.11	$\pm 0.03$	15.16 <sup>ab</sup>	$\pm 4.73$
<i>Fagus sylvatica</i>	0.35	$\pm 0.11$	6	32.45	0.05	$\pm 0.02$	6.60 <sup>b</sup>	$\pm 3.76$
<i>Betula pendula</i>	0.74	$\pm 0.08$	2	31.99	0.06	$\pm 0.02$	14.05 <sup>ab</sup>	$\pm 2.42$
<i>Castanea sativa</i>	0.41	$\pm 0.10$	4	33.62	0.17	$\pm 0.06$	1.05 <sup>b</sup>	$\pm 0.40$
<i>Quercus robur</i>	0.38	$\pm 0.10$	5	30.29	0.08	$\pm 0.03$	5.23 <sup>b</sup>	$\pm 1.23$
<i>Acer</i> <i>pseudoplatanus</i>	0.52	$\pm 0.11$	3	25.51	0.03	$\pm 0.02$	103.9 <sup>a</sup>	$\pm 54.77$
<b>0.2–0.3 m</b>								
<i>Alnus glutinosa</i>	0.29 <sup>ab</sup>	$\pm 0.05$	5	20.66	0.11	$\pm 0.03$	3.82	$\pm 1.95$
<i>Fraxinus excelsior</i>	0.90 <sup>a</sup>	$\pm 0.32$	1	17.43	0.11	$\pm 0.06$	10.67	$\pm 2.73$
<i>Fagus sylvatica</i>	0.17 <sup>b</sup>	$\pm 0.04$	7	16.18	0.02	$\pm 0.00$	8.96	$\pm 2.80$
<i>Betula pendula</i>	0.44 <sup>ab</sup>	$\pm 0.06$	3	19.30	0.08	$\pm 0.01$	6.95	$\pm 1.86$
<i>Castanea sativa</i>	0.36 <sup>ab</sup>	$\pm 0.13$	4	29.47	0.05	$\pm 0.02$	2.57	$\pm 1.05$

<i>Quercus robur</i>	0.26 <sup>ab</sup>	± 0.04	6	20.61	0.09	± 0.03	3.78	± 1.05
<i>Acer</i>								
<i>pseudoplatanus</i>	0.64 <sup>a</sup>	± 0.19	2	31.61	0.12	± 0.05	12.61	± 6.33
<b>0–0.3 m</b>								
<i>Alnus glutinosa</i>	1.39 <sup>b</sup>	± 0.26	4	N/A	0.51	± 0.11	3.43	± 1.20
<i>Fraxinus excelsior</i>	5.16 <sup>a</sup>	± 0.71	1	N/A	0.58	± 0.17	12.01	± 3.83
<i>Fagus sylvatica</i>	1.07 <sup>b</sup>	± 0.26	7	N/A	0.11	± 0.03	10.68	± 2.84
<i>Betula pendula</i>	2.30 <sup>ab</sup>	± 0.12	2	N/A	0.27	± 0.05	9.30	± 1.78
<i>Castanea sativa</i>	1.21 <sup>b</sup>	± 0.24	6	N/A	0.29	± 0.10	1.97	± 0.73
<i>Quercus robur</i>	1.26 <sup>b</sup>	± 0.34	5	N/A	0.30	± 0.09	4.57	± 0.81
<i>Acer</i>								
<i>pseudoplatanus</i>	2.04 <sup>ab</sup>	± 0.56	3	N/A	0.23	± 0.07	14.82	± 6.65

630

631 Table 4 Between-subject effects of species and depth on fine root traits including specific root length  
632 (m g<sup>-1</sup>), root area index (m<sup>2</sup> m<sup>-2</sup>), specific root surface area (m<sup>2</sup> kg<sup>-1</sup>), root length density (cm cm<sup>-3</sup>) and  
633 root tip density (× 10<sup>3</sup> m<sup>-2</sup>).

		df	F-statistic	p
Species	Specific root length (m g <sup>-1</sup> )	6	21.825	<0.001
	Root area index (m <sup>2</sup> m <sup>-2</sup> )	6	22.757	<0.001
	Specific root surface area (m <sup>2</sup> kg <sup>-1</sup> )	6	19.549	<0.001
	Root length density (cm cm <sup>-3</sup> )	6	29.519	<0.001
	Root tip density (× 10 <sup>2</sup> m <sup>-2</sup> )	6	25.816	<0.001
Depth	Specific root length (m g <sup>-1</sup> )	2	76.572	<0.001
	Root area index (m <sup>2</sup> m <sup>-2</sup> )	2	22.994	<0.001
	Specific root surface area (m <sup>2</sup> kg <sup>-1</sup> )	2	0.591	0.557
	Root length density (cm cm <sup>-3</sup> )	2	39.089	<0.001
	Root tip density (× 10 <sup>2</sup> m <sup>-2</sup> )	2	57.046	<0.001
Species×Depth	Specific root length (m g <sup>-1</sup> )	12	0.395	0.961
	Root area index (m <sup>2</sup> m <sup>-2</sup> )	12	0.826	0.623
	Specific root surface area (m <sup>2</sup> kg <sup>-1</sup> )	12	0.392	0.962
	Root length density (cm cm <sup>-3</sup> )	12	0.852	0.598
	Root tip density (× 10 <sup>2</sup> m <sup>-2</sup> )	12	1.108	0.370

634

635 **Table 5.** Fine root metrics (specific root length (m g<sup>-1</sup>), root area index (m<sup>2</sup> m<sup>-2</sup>), specific root surface  
636 area (m<sup>2</sup> kg<sup>-1</sup>), root length density (cm cm<sup>-3</sup>) and root tip density (× 10<sup>2</sup> m<sup>-2</sup>) of seven tree species (*Alnus*  
637 *glutinosa*, *Fraxinus excelsior*, *Fagus sylvatica*, *Betula pendula*, *Castanea sativa*, *Quercus robur* and *Acer*

638 *pseudoplatanus*) at three soil depths (0-0.1, 0.1-0.2, 0.2-0.3 m). SE =  $\pm 1$  standard error. Superscript letters denote Tukey post hoc comparisons ( $p < 0.05$ ) of  
639 root traits within each soil depth.

Root trait	Depth	<i>Alnus glutinosa</i>			<i>Fraxinus excelsior</i>			<i>Fagus sylvatica</i>			<i>Betula pendula</i>			<i>Castanea sativa</i>			<i>Quercus robur</i>			<i>Acer pseudoplatanus</i>		
Specific root length (m g <sup>-1</sup> )	0-0.1	8.27 <sup>b</sup>	$\pm$	1.94	22.23 <sup>a</sup>	$\pm$	3.62	26.38 <sup>a</sup>	$\pm$	3.22	14.88 <sup>ab</sup>	$\pm$	1.58	13.43 <sup>ab</sup>	$\pm$	1.22	20.20 <sup>a</sup>	$\pm$	3.04	22.43 <sup>a</sup>	$\pm$	3.57
	0.1-0.2	4.13 <sup>c</sup>	$\pm$	0.41	10.09 <sup>ab</sup>	$\pm$	0.45	13.45 <sup>a</sup>	$\pm$	3.85	6.36 <sup>abc</sup>	$\pm$	0.29	5.79 <sup>bc</sup>	$\pm$	0.83	9.88 <sup>ab</sup>	$\pm$	2.50	9.42 <sup>ab</sup>	$\pm$	0.38
	0.2-0.3	3.37 <sup>c</sup>	$\pm$	0.45	8.98 <sup>ab</sup>	$\pm$	0.73	11.78 <sup>a</sup>	$\pm$	1.66	5.35 <sup>bc</sup>	$\pm$	0.57	5.53 <sup>bc</sup>	$\pm$	0.95	9.31 <sup>ab</sup>	$\pm$	1.06	6.52 <sup>ab</sup>	$\pm$	1.12
Root area index (m <sup>2</sup> m <sup>-2</sup> )	0-0.1	0.88 <sup>b</sup>	$\pm$	0.27	6.02 <sup>a</sup>	$\pm$	0.86	0.96 <sup>b</sup>	$\pm$	0.26	1.27 <sup>b</sup>	$\pm$	0.19	0.51 <sup>b</sup>	$\pm$	0.09	0.78 <sup>b</sup>	$\pm$	0.26	1.25 <sup>b</sup>	$\pm$	0.32
	0.1-0.2	0.28 <sup>b</sup>	$\pm$	0.07	2.59 <sup>a</sup>	$\pm$	0.64	0.50 <sup>b</sup>	$\pm$	0.16	0.76 <sup>ab</sup>	$\pm$	0.11	0.47 <sup>b</sup>	$\pm$	0.10	0.45 <sup>b</sup>	$\pm$	0.09	0.70 <sup>b</sup>	$\pm$	0.17
	0.2-0.3	0.29 <sup>b</sup>	$\pm$	0.05	1.78 <sup>a</sup>	$\pm$	0.57	0.26 <sup>b</sup>	$\pm$	0.05	0.42 <sup>b</sup>	$\pm$	0.06	0.39 <sup>b</sup>	$\pm$	0.12	0.37 <sup>b</sup>	$\pm$	0.05	0.74 <sup>ab</sup>	$\pm$	0.20
Specific root surface area (m <sup>2</sup> kg <sup>-1</sup> )	0-0.1	10.29 <sup>c</sup>	$\pm$	1.87	19.40 <sup>a</sup>	$\pm$	0.41	16.97 <sup>ab</sup>	$\pm$	1.73	11.14 <sup>bc</sup>	$\pm$	0.78	11.58 <sup>bc</sup>	$\pm$	0.60	14.32 <sup>abc</sup>	$\pm$	1.36	14.90 <sup>abc</sup>	$\pm$	1.47
	0.1-0.2	10.80 <sup>b</sup>	$\pm$	0.68	21.13 <sup>a</sup>	$\pm$	0.89	15.60 <sup>ab</sup>	$\pm$	1.96	10.17 <sup>b</sup>	$\pm$	0.36	11.59 <sup>b</sup>	$\pm$	1.28	13.74 <sup>b</sup>	$\pm$	2.74	13.17 <sup>b</sup>	$\pm$	0.66
	0.2-0.3	9.95 <sup>cd</sup>	$\pm$	0.70	20.15 <sup>a</sup>	$\pm$	1.30	15.92 <sup>ab</sup>	$\pm$	1.69	9.50 <sup>d</sup>	$\pm$	0.60	11.68 <sup>bcd</sup>	$\pm$	0.94	14.15 <sup>abc</sup>	$\pm$	0.85	11.92 <sup>bcd</sup>	$\pm$	1.27
Root length density (cm cm <sup>-3</sup> )	0-0.1	0.70 <sup>b</sup>	$\pm$	0.24	6.56 <sup>a</sup>	$\pm$	0.65	1.51 <sup>b</sup>	$\pm$	0.42	1.70 <sup>b</sup>	$\pm$	0.33	0.57 <sup>b</sup>	$\pm$	0.07	1.03 <sup>b</sup>	$\pm$	0.27	1.82 <sup>b</sup>	$\pm$	0.43
	0.1-0.2	0.21 <sup>c</sup>	$\pm$	0.05	2.42 <sup>a</sup>	$\pm$	0.60	0.72 <sup>b</sup>	$\pm$	0.19	0.94 <sup>ab</sup>	$\pm$	0.15	0.43 <sup>bc</sup>	$\pm$	0.07	0.61 <sup>b</sup>	$\pm$	0.10	0.96 <sup>ab</sup>	$\pm$	0.20
	0.2-0.3	0.19 <sup>c</sup>	$\pm$	0.04	1.55 <sup>a</sup>	$\pm$	0.47	0.37 <sup>bc</sup>	$\pm$	0.06	0.46 <sup>bc</sup>	$\pm$	0.05	0.33 <sup>bc</sup>	$\pm$	0.07	0.48 <sup>bc</sup>	$\pm$	0.09	0.77 <sup>ab</sup>	$\pm$	0.19
Root tip density (x 10 <sup>2</sup> m <sup>-2</sup> )	0-0.1	242.89 <sup>bc</sup>	$\pm$	45.28	1275.01 <sup>a</sup>	$\pm$	199.30	515.07 <sup>ab</sup> <sub>c</sub>	$\pm$	127.84	625.53 <sub>ab</sub>	$\pm$	150.04	174.35 <sup>c</sup>	$\pm$	17.17	348.08 <sup>bc</sup>	$\pm$	88.43	528.29 <sup>ab</sup>	$\pm$	116.41
	0.1-0.2	63.31 <sup>c</sup>	$\pm$	11.36	512.43 <sup>a</sup>	$\pm$	132.38	260.47 <sup>ab</sup>	$\pm$	64.56	307.72 <sub>a</sub>	$\pm$	23.23	104.35 <sup>bc</sup>	$\pm$	14.62	205.66 <sup>ab</sup>	$\pm$	39.54	246.19 <sup>ab</sup>	$\pm$	36.84
	0.2-0.3	59.83 <sup>c</sup>	$\pm$	9.99	314.58 <sup>a</sup>	$\pm$	84.47	130.93 <sup>ab</sup> <sub>c</sub>	$\pm$	17.09	137.32 <sub>ab</sub>	$\pm$	12.12	85.35 <sup>bc</sup>	$\pm$	16.00	157.04 <sup>ab</sup>	$\pm$	25.09	205.31 <sup>a</sup>	$\pm$	38.81

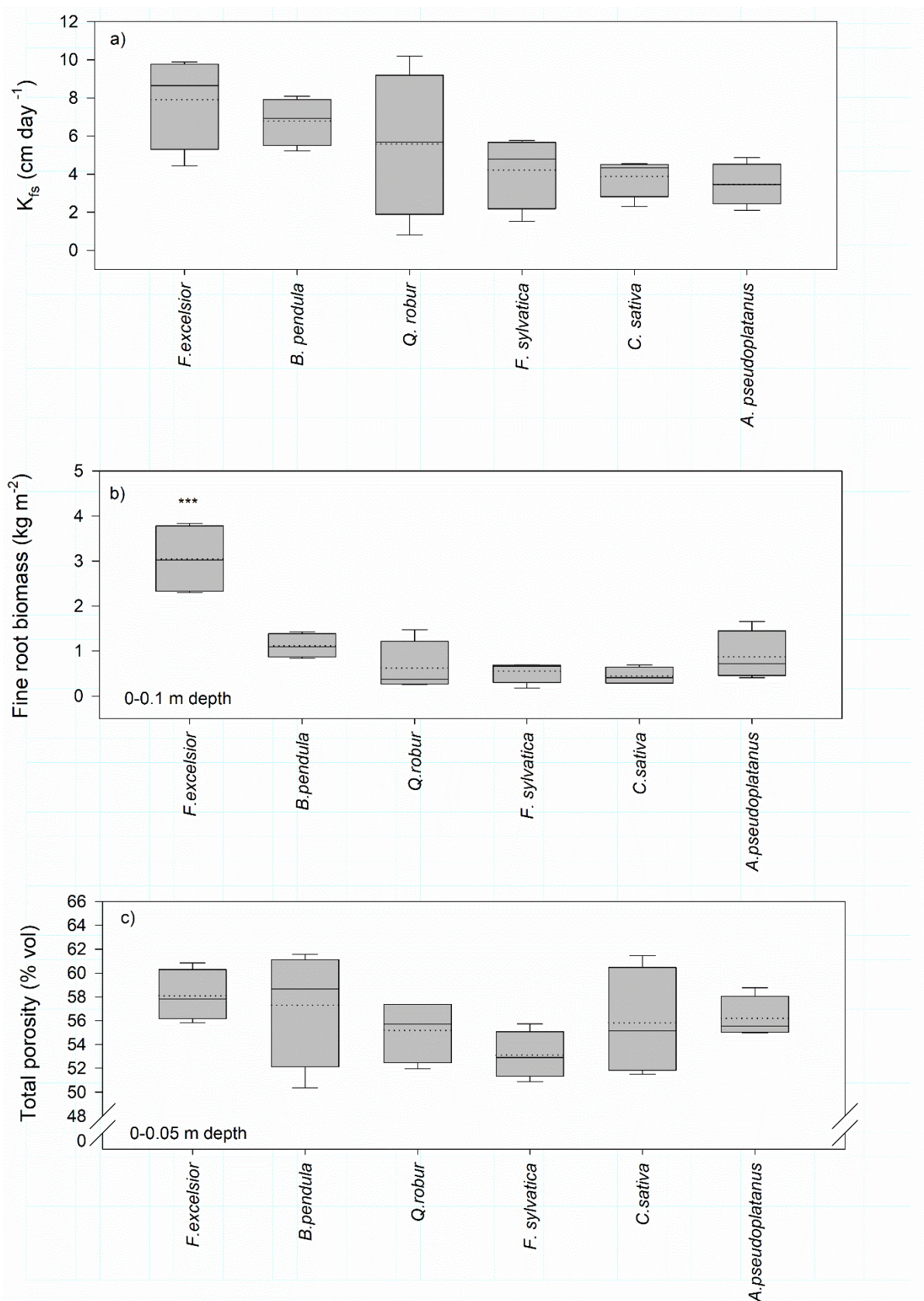
640

641

642 **Table 6.** Main effects of four contrasting soil textures' (Rendzic Leptosol, silty-clay loam – limestone  
643 rich; Haplic Luvisol, silty-clay loam; Dystric Fluvisol Cambisol, sandy silt loam; Dystric Gleysol, clay loam)  
644 fine root biomass in three soil depths (0-0.1, 0.1-0.2, 0.2-0.3 m).

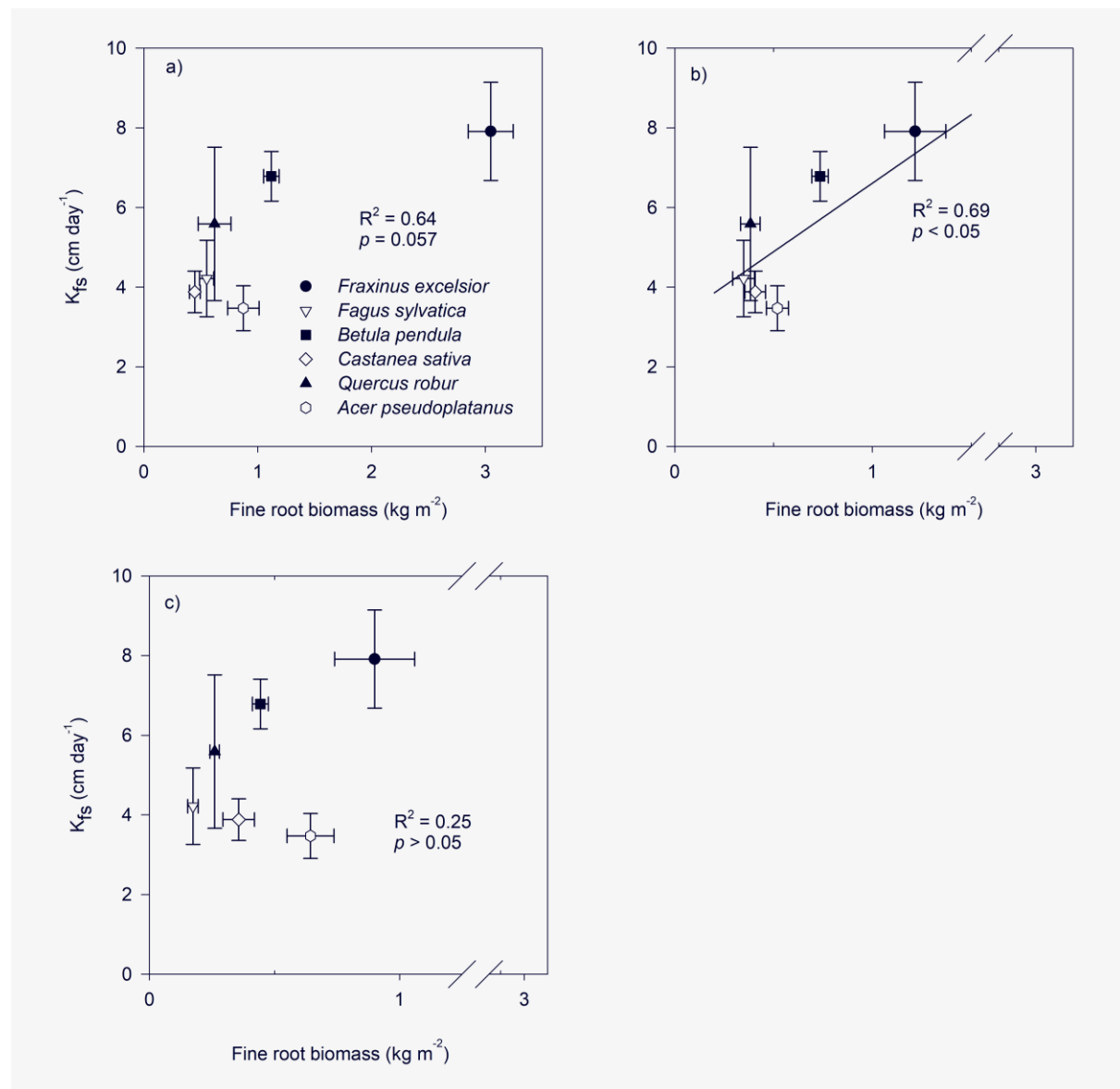
Factor	df	F	<i>p</i> -value
Depth	2	24.107	0.000
Soil class	3	6.394	0.002
Depth*Class	6	1.185	0.347

645 **Figure legends**



**Fig. 1.** Variation in soil and fine root properties amongst plots ( $n=4$ ) of six tree species: (a) Surface field saturated hydraulic conductivity ( $K_{fs}$ ;  $\text{cm day}^{-1}$ ), (b) fine root biomass ( $\text{kg m}^{-2}$ ) in the 0-0.1 m depth, (c)

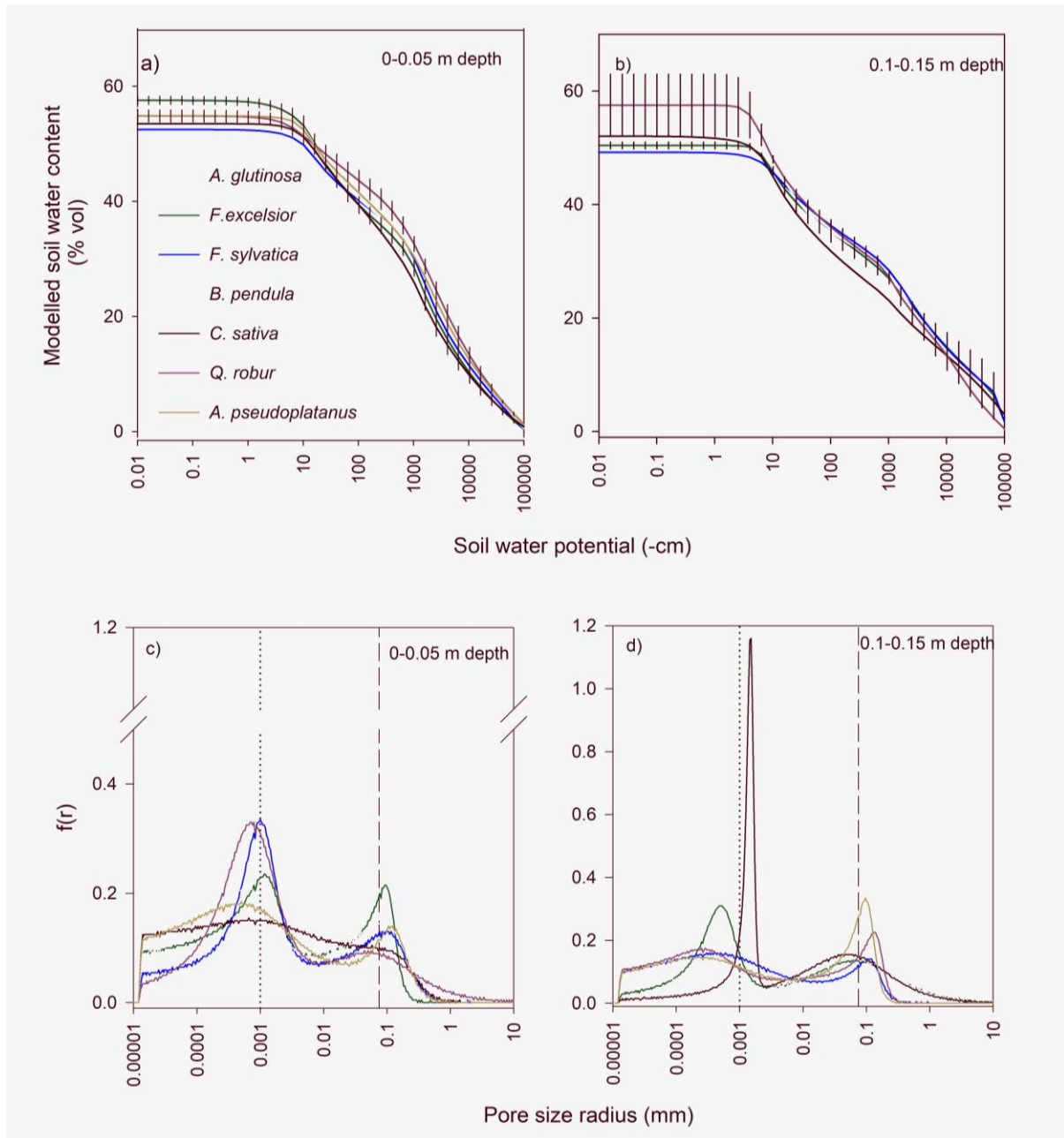
total soil porosity (% volume) calculated from cores (excluding stone fraction) taken from the 0-0.05 m depth. Data shown are mean (dashed horizontal line) and median (solid horizontal line). The boxes define quartiles and whiskers  $\pm$  one standard error. For fine root biomass, there was a species main effect  $p < 0.001$ . No statistically significant differences were found in  $K_{fs}$  or total porosity amongst species ( $p > 0.05$ ). *Alnus glutinosa* is excluded from biomass analysis because the stand was in poor health, demonstrated by a large fraction of necromass amongst the fine roots.



**Fig. 2.** Relationship between mean plot (n=4) surface field-saturated hydraulic conductivity ( $K_{fs}$ ; cm day<sup>-1</sup>) and fine root biomass (kg m<sup>-2</sup>) for six species, *F. excelsior*, *F. sylvatica*, *B. pendula*, *C. sativa*, *Q.*

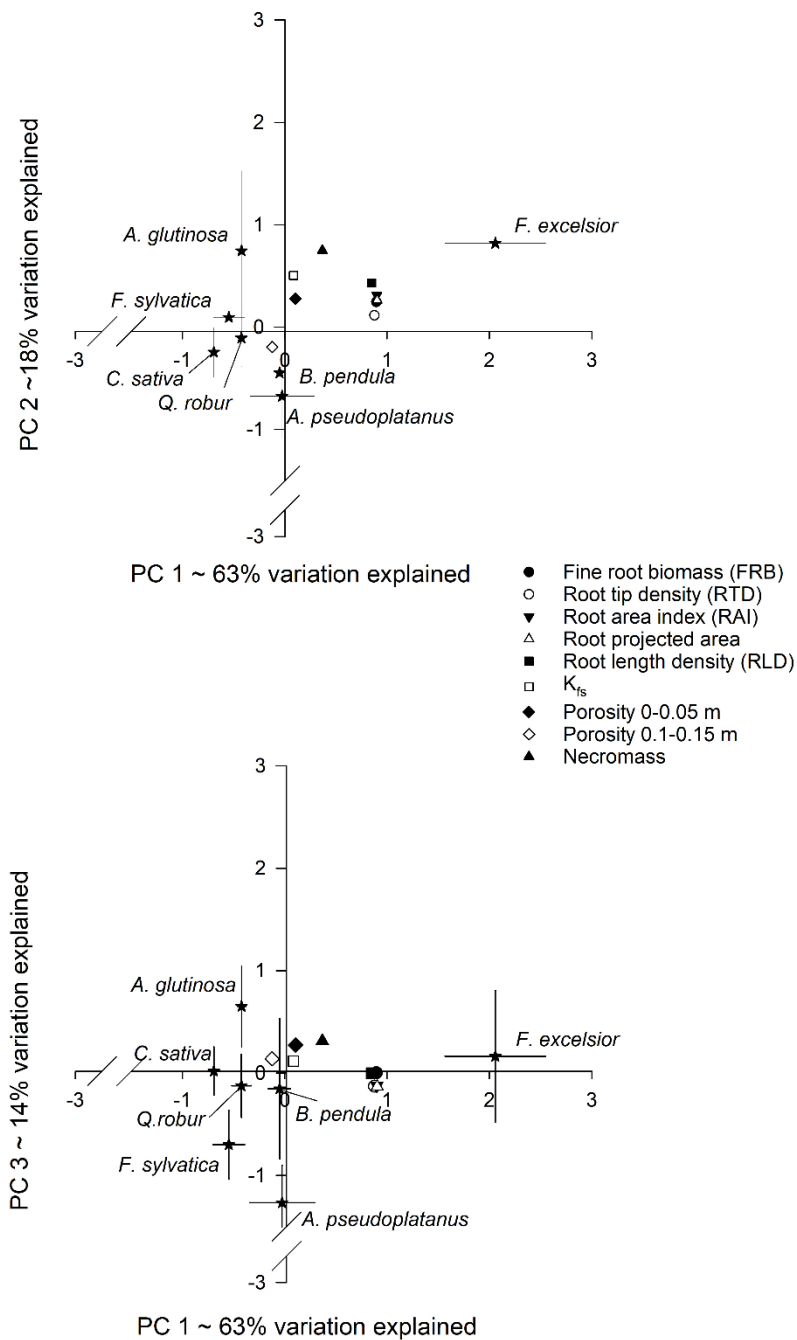


*robur* and *A. pseudoplatanus*, in the (a) 0-0.1 m, (b) 0.1-0.2 m and (c) 0.2-0.3 m soil depths. Data shown are mean  $\pm$  one standard error for each species. *Alnus glutinosa* is excluded from biomass analysis because the stand was in poor health, demonstrated by a large fraction of necromass amongst the fine roots.

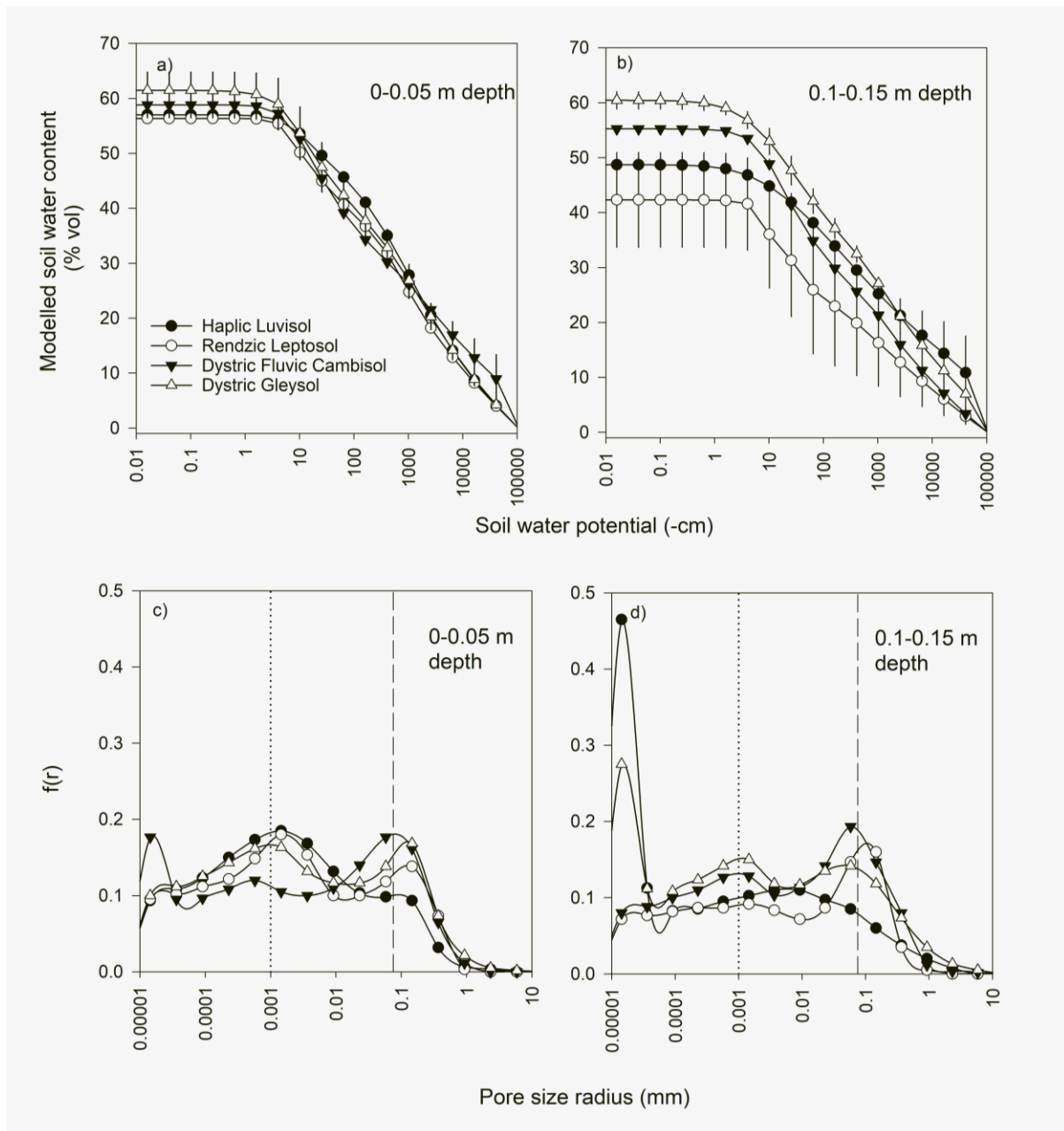


**Fig. 3.** Mean soil water retention curves for plots (n=4) of seven tree species in the (a) 0-0.05 m and (b) 0.01-0.15 m depths. The data are modelled using the bimodal Fredlund-Xing PDI model using

666 measured soil water content and potential (HYPROP) data. Modelled effective pore-size radius  
667 distribution (Blonquist et al., 2006), displayed on a common log scale, of the seven species in the (c)  
668 0-0.05 m and (d) 0.1-0.15 m depths. The pore-size distribution ( $f(r)$ ) represents the proportional  
669 volume of the combined effective pore size radii. Values to the right of the dotted vertical line indicate  
670 pore radius sizes where capillary forces dominate water movement (Kosugi et al., 2002). Values to the  
671 right of the dashed vertical line indicate macropore radius sizes  $> 0.075$  mm.

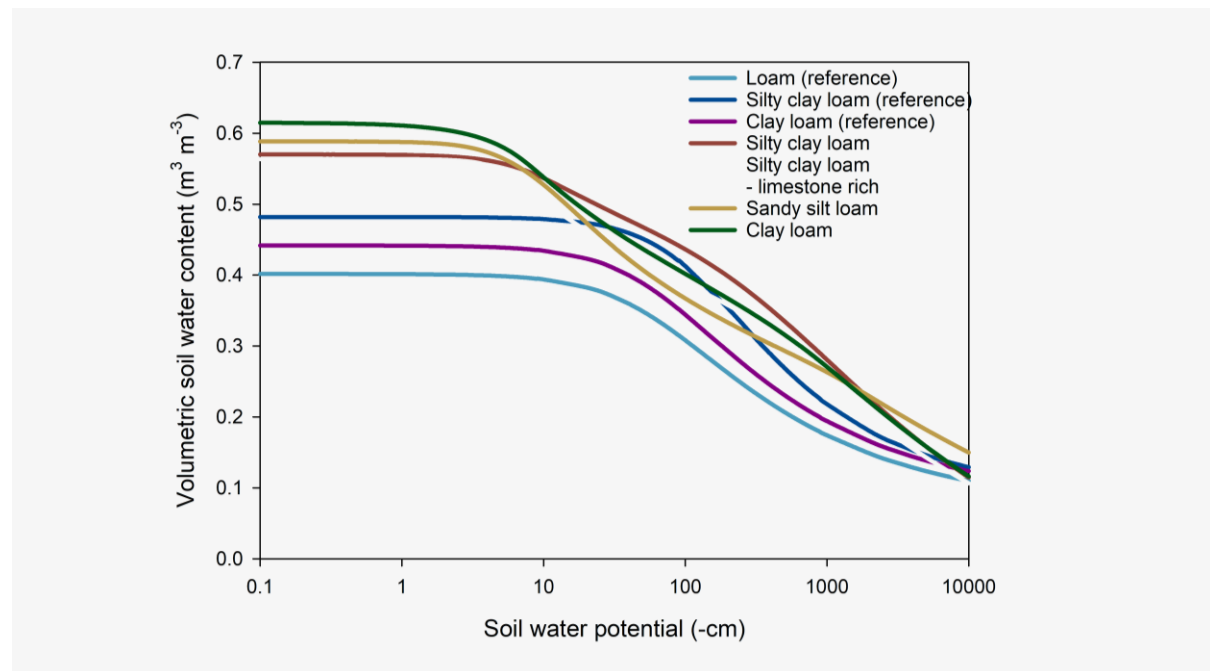


**Fig. 4.** Principal component analysis examining the relationships between field saturated hydraulic conductivity ( $K_{fs}$ ), tree species (*A. glutinosa*, *F. excelsior*, *B. pendula*, *F. sylvatica*, *C. sativa*, *Q. robur* and *A. pseudoplatanus*), fine root morphological variables (root biomass, root tip number, root area index, root projected area, root length density and necromass) and soil porosity (% volume) at two soil depths (0-0.05 and 0.1-0.15 m). Error bars represent  $\pm$  one standard error.



**Fig. 5.** Mean soil water retention curves for four sites with contrasting soil classes: Haplic Luvisol (silty clay loam); Rendzic Leptosol (silty clay loam - limestone rich); Dystric Fluvic Cambisol (sandy silt loam); and Dystric Gleysol (clay loam), at (a) 0-0.05 m and (b) 0.01-0.15 m depths. The data are modelled using the bimodal Fredlund-Xing PDI model (Fredlund and Xing, 1994) using measured soil water content (HYPROP) data. Modelled pore-size distribution (Blonquist et al., 2006) displayed on a common log scale from contrasting soil classes at (c) 0-0.05 m and (d) 0.1-0.15 m depths. The pore-size distribution ( $f(r)$ ) represents the proportional volume of the combined effective pore size radii.

Values to the right of the dotted vertical line indicate pore sizes where capillary forces dominate water movement (Kosugi et al., 2002). Values to the right of the dashed vertical line indicate macropore pore sizes  $> 0.075$  mm. Values between the vertical lines indicate mesopores.



**Fig. 6.** Soil water retention curves (SWRC) for the four soil textures used in our study: Haplic Luvisol (silty clay loam); Rendzic Leptosol (silty clay loam - limestone rich); Dystric Fluvisol Cambisol (sandy silt loam); and Dystric Gleysol (clay loam), and modelled SWRC for three reference soil textures (loam, silty clay loam and clay loam) from the Rosetta modelling framework for pedotransfer functions (Schaap et al., 2001).  $pF$ , the decimal log of soil water potential (cm), describes the amount of force or suction required to extract water from the soil.