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# Describing the vertical root distribution of alpine plants with simple climate, soil, and plant attributes

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- 15 Abstract

16 The vertical root distribution (VRD) in the soil remains unknown for most plant species, as 17 studying root systems in different pedo-climatic settings is time-consuming and 18 methodologically challenging. Yet, information on the VRD of different vegetation types is 19 essential to understand better the biogeochemical processes occurring at the soil-plant-20 atmosphere continuum. The aim of this study was to describe the (VRD) of three dominant 21 alpine, herbaceous plants (i.e. Euphrasia minima Jacq., Leucanthemopsis alpina L., and Poa 22 alpina L.) on the basis of simple and easy-to-measure climate, soil, and plant attributes in order to test the validity of existing descriptive protocols and parametric ecohydrological models. 23 24 The results showed that the VRD decreased with soil depth for the three plants and that it can 25 be effectively described with a negative exponential equation. Key VRD parameters, such as

the mean rooting depth, cross-sectional area at the root collar, and root biomass, were both site and species-specific but they were chiefly influenced by the attributes regulating the soil's water mass balance. The existing parametric ecohydrological models were not able to portray successfully the VRD of the studied alpine plants but we found a strong correlation between empirical and parametric VRD models that establish a clear direction for future research. Future work should address the influence of the snowpack characteristics and the length of the snow-free and frozen ground periods on the soil's ecohydrology and VRD in alpine ecosystems.

- 35 Keywords: root, model, ecohydrological, alpine, data mining

#### 37 Abbreviations

	M	00	
α	Mean precipitation intensity over VSD	θfc	Soil moisture at field capacity
AI	Aridity index	$\theta g$	Gravimetric soil moisture content
ALR	Allometry ratio	$\theta wp$	Soil moisture at wilting point
Ar	Root cross-sectional area	$ ho_{bk}$	Soil's bulk density
Aro	Cross-sectional area at root collar	$ ho_r$	Root mass density
b	Mean rooting depth	Sa	Soil's sand content
Cl	Soil's clay content	Sk	Soil skeleton
CN	Concave topographic curvature	SOC	Soil organic carbon
CS	Plant's crown spread	Sp	Plant's aerial projected area
CX	Convex topographic curvature	Tbase	Optimum temperature for plant growth
Etp	Potential evapotranspiration	Tmn	Daily minimum air temperature
FL	Flat topographic curvature	Tmx	Daily maximum air temperature
GDD	Growing-degree day	VRD	Vertical root distribution
λ	Precipitation frequency over VSD	VSD	Vegetative season duration
Ma	Aboveground biomass	WAP	Water available to plants
Mr	Belowground biomass	Z	Soil depth
n	Soil porosity		

46

47 1. Introduction

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49 Knowledge of root systems of different vegetation types is essential for a better understanding 50 of biogeochemical processes occurring at the soil-plant-atmosphere continuum (Rodriguez-51 Iturbe and Porporato, 2005). Despite relatively recent efforts in investigation and description 52 of plant root architecture (e.g., Waisel et al., 2002, Mickovski and Ennos 2003, Mickovski and 53 van Beek 2005), it remains largely unexplored for most plant species. A number of authors 54 have explored root system architecture of a large number of plants native to almost all 55 bioclimatic regions (e.g. Schenk and Jackson, 2005), and attempted prediction of rooting 56 depths through optimisation (van Wijk and Bouten, 2001; Kleidon, 2004) or inverse methods 57 (Zuo and Zhang, 2002). These approaches led to an increase in knowledge and understanding 58 of plant physiological processes such as water and nutrient uptake, resources competition, and 59 plant-soil interactions (Herbert et al., 2004; Laio et al., 2006, Preti et al., 2010; Gonzalez-60 Ollauri and Mickovski, 2017a). Despite the efforts in the past decade, the comprehensive 61 understanding of the effect of soil and plant properties and climate conditions on root 62 architecture and morphology remains largely unknown.

63

Obtaining root information is time-consuming and methodologically challenging. The investigation of root systems normally involves destructive and invasive sampling approaches (Bhöm, 1979), followed by detailed description and measurement of specific root traits (Stokes et al., 2009). However, for many environmental applications related to plant-soil interactions, knowledge of the vertical root distribution (VRD) – i.e. the pattern in which root density biomass is distributed along the soil profile - is perhaps the most important feature to know 70 because it can be used, for example, to estimate the degree of soil-root mechanical 71 reinforcement (Arnone et al., 2016; Gonzalez-Ollauri and Mickovski, 2016; Kokutse et al., 72 2016), to estimate plant-water uptake (Jarvis, 1989; Laio, 2006; Shukla, 2014), or to gain 73 insight into the ability of vegetation to remove pollutants from the soil (Verma et al., 2006; 74 Gonzalez-Ollauri and Mickovski, 2018; Lucherini et al., 2020). For most of these applications, 75 VRD can be easily portrayed with asymptotic mathematical functions (Jackson et al., 1996), which substantially simplify the process of describing the root system, as they normally require 76 77 few parameters related to the root system, such as the rooting depth or the cross-sectional area 78 at the root collar (Preti et al., 2010). However, standard and reproducible protocols to describe 79 VRD are still lacking.

80

81 The way in which roots distribute in the soil has been the scope of research for many decades 82 (e.g. Darwin, 1880; Laio et al., 2006). Previous research indicates that the root distribution in 83 the soil is mostly influenced by water availability to plants (i.e. hydrotropism; Tsutsumi et al., 84 2003). This is relevant because it permits to connect VRD to climate and soil attributes 85 regulating the water cycle in the soil (e.g. soil porosity, soil organic matter, soil texture, etc; 86 Toth et al., 2015), and to set the basis for establishing cost-effective analytical approaches 87 describing VRD on the basis of few, easy-to-measure variables. As a result, and to the best of 88 our knowledge, two parametric, ecohydrological models predicting VRD have been developed 89 for arid or semi-arid (Laio et al., 2006; Preti et al., 2010) and for temperate-humid (Gonzalez-90 Ollauri and Mickovski, 2016) ecosystems, respectively. These models incorporate plant-91 specific attributes by considering the relative allocation of plant biomass between the above-92 and belowground soil compartments (i.e. allometry; Cheng and Niklas. 2007) and by assuming 93 that the root system can take a known, simple geometrical shape, such as a cylinder, a cone, or 94 a hemisphere (e.g. Lynch, 1995; Kutschera and Lichtenegger, 1992; Kokutse et al., 2006). The

95 ability of these models to realistically portray VRD has been successfully tested for few herbaceous (Gonzalez-Ollauri and Mickovski, 2016) and shrub species (Preti and Giadrossich, 96 97 2009; Preti et al., 2010), but their application to the wider pool of plants, in general, and to 98 woody plants (e.g. Preti et al., 2010; Tron et al., 2014; Tardio et al., 2016), in particular, needs 99 further validation. In addition, the robustness of the model conceptualisation and the 100 assumptions still need to be verified against primary data showing the influence of multiple 101 soil and plant attributes on VRD, which could help to identify potential model improvements 102 including application to climates with different ecohydrological features, such as tropical or 103 alpine.

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105 Alpine ecosystems are normally found above the upper limit of tree growth in mountainous 106 areas. They are generally characterised by cold winter temperatures, precipitation in the form 107 of snow, and short snow-free periods (Freppaz et al., 2019) - all of which tend to limit the 108 duration of the vegetative season. As a result, alpine vegetation, which mostly comprises low-109 growing herbaceous perennial plants, tends to be sparse and endemic to these ecosystems, or 110 it may have evolved to withstand the environmental stress related to alpine climates (Germino, 111 2014). In addition, the growth and development of alpine plants is also constrained by the low 112 availability of nutrients in the soil, particularly nitrogen (e.g. Freppaz et al., 2019; Zong et al., 113 2020). Still, alpine vegetation may play a crucial role in cycling carbon and nutrients in alpine 114 ecosystems (e.g. Iversen et al., 2014), or in protecting the soil against shallow landslides and 115 erosion (Preti. 2013; Burylo et al., 2014; Gonzalez-Ollauri and Mickovski, 2017b), where 116 poorly developed soils subjected to freezing are prone to soil mass wasting processes (e.g. Hudek et al., 2017a). Yet, knowledge of the root systems in alpine plants is scarce (e.g. Iversen 117 118 et al., 2014) and only few studies have attempted addressing this knowledge gap (e.g. Pohl et 119 al., 2011; Burylo et al., 2014; Hudek et al., 2017b).

121 The aim of this study is to describe the VRD of three dominant alpine plants on the basis of simple and easy-to-measure climate, soil, and plant attributes in order to test existing 122 123 descriptive protocols and parametric ecohydrological models. The objectives of the study 124 include (i) selection and characterisation of three alpine sites in terms of climate and soil 125 attributes, (ii) sampling above- and below-ground plant parts of three dominant pioneer plant species from the three study locations to retrieve relevant plant information and to describe the 126 127 VRD, (iii) investigation of the influence of evaluated soil and plant attributes on key VRD 128 parameters, and (iv) testing the predictive capacity of an existing parametric ecohydrological 129 model for VRD using pedo-climatic and plant attributes.

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131 2. Materials and Methods

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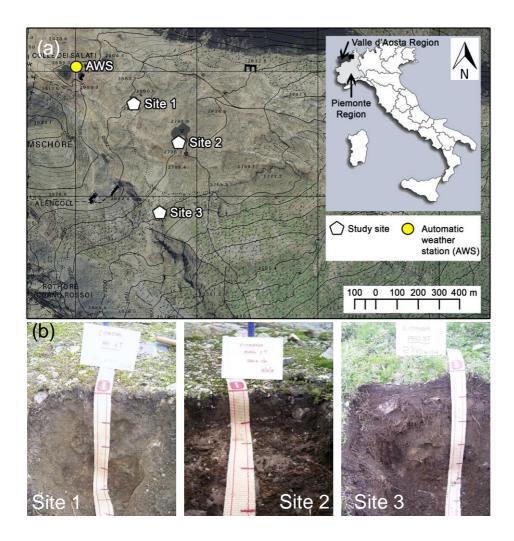
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135 The study area is located adjacent to Monte Rosa massif (4634 m above sea level; a.s.l.), within the Long-Term Ecological Research (LTER) site Angelo Mosso Institute, Valle d'Aosta 136 137 Region, Northwest Italy (Fig. 1a). The climate in the study area is Alpine (ET; Köppen, 1884), 138 characterised by a mean annual air temperature of -2.5°C, a cumulative annual snowfall of 8.50 139 m, and a mean annual rainfall of 350 mm. The snowpack generally develops in late October -140 early November and lasts until the onset of snowmelt in late May – early June. Soil temperature 141 and meteorological parameters such as air temperature, rainfall during the snow-free season, 142 and snowfall have been continuously recorded with a 1-minute resolution in the study area 143 since 2005 using one Automatic Weather Station (AWS) located at 2901 m a.s.l. (Fig. 1; 144 Comando Truppe Alpine - Servizio Meteomont).

<sup>133 2.1.</sup> Study area

146 Three, high-altitude study sites located in the upper part of a glacier valley were selected within 147 the study area (Fig. 1a). The sites were located at different elevations and had distinct 148 underlaying soil types -i.e. Site 1: 2840 m a.s.l on Dystric Leptic Regosol; Site 2: 2795 m on 149 Dystric Lithic Leptosol ; Site 3: 2770 m on Haplic Umbrisol (IUSS Working Group WRB, 150 2015). Sites 1 and 2 (Figs.1b and 1c) are characterised by relatively flat topographies. The 151 dominant vegetation at these sites comprises nival plant species of perennial grasses (Poa laxa, 152 Poa alpina), together with other herbaceous species such as forbs (Leucanthemopsis alpina, 153 Gnaphalium supinum), cushion plants (Minuartia sedoides) and dwarf, woody plants that can 154 tolerate long-lasting snow cover (Salix herbacea, Loiseleuria procumbens). The topography of 155 Site 3 is rougher and the vegetation cover denser than at sites 1 and 2 (Fig. 1a). Alpine 156 grasslands dominate Site 3 with the most characteristic plant species being Carex curvula and 157 Euphorbia minima (Freppaz et al., 2019; Lonati et al., unpublished data).

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160

161 Figure 1 a) Locations of the study sites (1-3) and b) characteristic soil profiles for site 1, 2 and 3.

163 2.2.Climatic attributes

We measured four climatic attributes reported to influence plant growth and root development (e.g. Preti et al., 2010; Gonzalez-Ollauri and Mickovski, 2016), and which are nested in the parametric, ecohydrological VRD model to be tested herein (Section 2.6; Table 1). To this end, we examined daily meteorological records collected with one AWS between 2005 and 2019 (Fig. 1a), which we assumed to be representative for the three study sites. The vegetative season duration (VSD) was defined using the heuristic growing-degree days (GDD; Eq.8) approach (e.g. McMaster and Wilhelm, 1997), which is a measure of the daily heat accumulation to

172 predict plant development and phenology. We assumed that the vegetative season begun once the cumulative GDD reached 200°C (Eq.8; Table 1), and that it ended when daily mean soil 173 174 temperature was below 4°C (i.e. root growth is inhibited under 4°C; e.g. Alvarez-Uria and 175 Körner, 2007). We also assumed 5°C as the optimum soil temperature for plant growth (*Tbase*; Eq.8; Table 1). We then calculated the aridity index (AI) of the study site as the ratio of the 176 177 total potential evapotranspiration to the total precipitation (Eq. 9; Table 1; Greve et al., 2019) 178 over the vegetative season. The total precipitation was considered as the sum of rainfall and 179 snowfall (i.e. snow water equivalent) recorded during the vegetative season of each examined 180 year. The total potential evapotranspiration over the vegetative season, which is in turn nested 181 in the VRD model (Eq.4; Table 1), was calculated with the Priestly and Taylor (1972) equation 182 (Eq. 10; Table 1). Subsequently, we estimated the mean precipitation depth ( $\alpha$ ; mm event<sup>-1</sup>) 183 and the frequency of precipitation events ( $\lambda$ ) during the vegetative season (Laio et al., 2006; 184 Preti et al., 2010). We calculated  $\alpha$  as the ratio of the total precipitation to the number of 185 precipitation events (i.e. days with precipitation > 0.2 mm) during the vegetative season 186 averaged for the studied time period comprised between 2005 and 2019. Rainfall lost to surface 187 runoff was assumed to be negligible in our study area (e.g. Tron et al., 2014). We calculated  $\lambda$ 188 as the ratio of the number of precipitation events to the total vegetative season duration 189 averaged for the studied period.

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191 2.3.Soil attributes

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The slope gradient and aspect were measured manually with a hand-held inclinometer and with a compass, respectively, at locations adjacent to each sampled plant individual (Section 2.4). The terrain curvature of each sampling location was visually described as either concave (CN), convex (CX) or flat (FL) (e.g. Gonzalez-Ollauri and Mickovski, 2017c). Undisturbed soil 197 samples (N=36) from the topsoil (0 mm - 100 mm below ground level; b.g.l) were collected at 198 the same locations where plants were sampled using a soil core sampler. These samples were 199 used to determine the soil bulk density ( $\rho_{bk}$ ; g cm<sup>-3</sup>), soil porosity (*n*) and gravimetric moisture content ( $\theta_g$ ; %) following standard methods (Head, 1980). The soil organic carbon (SOC; %) 200 201 and *pH* were determined using a portion of the collected soil materials, which was air-dried for 202 168 h and sieved through a 2 mm opening sieve. SOC was determined using a C/H/N analyser (Elementar Vario EL) while soil pH was determined in soil-water suspension (soil: water = 203 204 1:2.5) following the slurry method (ASTM, 1995) and using a pH electrode (Fisher Scientific 205 Accumet Basic AB15). Additional soil materials in form of bulk samples of 4kg-5kg were 206 collected with a shovel from the topsoil (0 mm – 300 mm b.g.l) at three representative locations 207 per study site (N=9). These representative sampling locations, which were assumed to capture 208 the main soil features within a study site, were within the area range covered by plant sampling, 209 and were less than 5 m away from any given plant individual sampled in this study. The soil 210 samples were stored in heavy-duty PVC bags and transported to the laboratory where they were 211 mixed per study site (N=3) prior to further analysis. The particle size distribution (PSD) of the 212 collected soil materials was determined through the dry sieving and the hydrometer methods 213 for the coarse (i.e. gravel and sand) and fines (i.e. silt and clay) fractions, respectively (Head, 214 1980). The soil skeleton (i.e. percentage of rock fragments in the soil sample; Sk; %) was 215 determined through dry sieving (Head, 1980). Soil moisture content at field capacity ( $\theta fc$ ; %) 216 and wilting point ( $\theta wp$ ; %) were estimated through pedotransfer functions (Eqs. 11 and 12; 217 Table 1; Toth et al., 2015), which are nested in the VRD model and use PSD, n, and SOC as 218 inputs.

219

220 2.4. Plant species, plant attributes and vertical root distribution

We selected three dominant, characteristic plant species for the study from sites 1, 2, and 3 (Fig.1):

- (i) dwarf eyebright (*Euphrasia minima* Jacq.), an annual, facultative root hemiparasite
  (Matthies, 1998; Fig. 2a) with erect stems reaching up to 150 mm, which grows in
  humid mountainous habitats between 950-3000 m a.s.l. (Asturnauta, 2020)
- (ii) alpine chrysanthemum (*Leucanthemopsis alpina* L.), a perennial, herbaceous plant
  belonging to the daisy family and specific to high alpine elevations, growing
  between 1800-3500 m a.s.l. It can be an early coloniser after the retreat of glaciers,
  being generally small in size (< 200 mm in height) with a root system characterised</li>
  by horizontal rhizomes (ukwildflowers.com, 2020; Fig. 2b).
- (iii) alpine bluegrass (*Poa alpina* L.), a subartic-alpine meadow tufted grass found in moist to dry limestone and in basaltic rock crevices and exposed heathlands. It is a pseudoviviparous, apomictic, and fast germinating plant (Pierce et al., 2000) that can reach up to 400 mm in height, normally has narrow leaves (2-4 mm), and its inflorescence is pyramidal, twice as tall as wide; it also has an adventitious root system that arises extra-vaginally through the lead-sheaths at the base of the plant (Pierce et al., 2000; Fig. 2c).



240 Figure 2. Selected root systems of (a) Euphrasia minima (b) Leucanthemopsis alpina (c) Poa alpina.

241 Plant sampling was undertaken at the height of the growing season when four individuals per 242 plant species were sampled at random locations within each study site (N=36; Fig. 1). The 243 projection area of the aerial plant parts (Sp; cm<sup>2</sup>; total plant aboveground area projected on the 244 ground, assuming a plant crown with a circular shape) and average crown spread (CS; cm; 245 mean spread diameter of the aboveground plant parts) were measured for each sampled 246 individual with a meter tape following the Spokes distance method (Blozan, 2006). Each plant 247 individual was excavated by hand before being clipped with scissors above the root collar (Fig. 248 3) to separate the above from the belowground part. The aboveground plant materials were 249 oven-dried at 60°C until constant mass to measure the aboveground biomass (Ma; g) of each 250 sampled individual. The belowground parts were cleaned with a water jet to separate soil 251 particles attached to the root system prior to being air-dried for 2 h for describing the vertical 252 root distribution (VRD).

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VRD was measured manually as the total cross-sectional root area at a given soil depth (Fig.
3) for each sampled plant before being averaged per plant species and study site. Using a
permanent marker and a ruler, marks were drawn on the root systems at equal length intervals

257 ranging from 5 to 20 mm and starting from the root collar (Fig. 3) to visualise the assumed 258 root-soil intersection planes (Fig. 3). The diameter of all the roots intersecting each plane was 259 measured with Vernier callipers, and their cross-sectional area  $(Ar(z); mm^2)$  was calculated 260 with Eqs. 1 and 2 (Table 1; Fig. 3), assuming all roots had circular cross-section when crossing 261 a given intersection plane (Fig. 3). Subsequently, Ar was averaged per intersection plane and 262 per plant species for a given sampling site. Then, a nonlinear least squares (nls) exponential model of the form  $y = ae^{-x/b}$ , where y is the dependent variable (i.e. root cross-sectional area; 263 Ar), x is the independent variable (i.e. soil depth), and a and b are fitting parameters (Eq.3; 264 265 Table 1) was fitted to the measured data resulting from the previous step. With this approach 266 and, for modelling purposes, it was assumed that the root biomass is distributed in the soil 267 following a cone shape volume (Fig. 3b; Preti et al., 2010; Gonzalez-Ollauri and Mickovski, 268 2016; see Supplementary Material) in which the total rooting depth (i.e. soil depth at which 95 269 % of roots are found; 3xb; mm; Laio et al., 2006) was the cone's height and the cross-sectional 270 area of the root collar (Aro; mm<sup>2</sup>) the cone's basal area (Fig. 3b). Accordingly, the rooting 271 depth (b; mm) was quantified as 1/3 of the longitudinal distance between the root collar and 272 the tip of the root system of each studied individual (Laio et al., 2006). It must be borne in 273 mind that with the former cone-shape-volume approach (Fig. 3b), we are not trying to capture 274 the shape of the root system per se (Fig. 2; e.g. Köstler et al., 1968) but to provide a generic 275 basis to model the widely-observed decrease in root biomass with soil depth (e.g. Schenk, 276 2005; see Supplementary Material). Finally, the root materials were oven-dried at 60°C until 277 constant mass to measure the root biomass (Mr; g) and the allometry ratio (ALR) as the quotient 278 between Mr and Ma.

280 Table 1. List of equations used in this study. Arid and humid ecosystems are defined on the basis of the aridity index (AI) over the growing season -i.e. AI < 1: arid; AI > 1: humid; ppu: parts-per

#### 281 unit. VS: vegetative season.

Definition	Equation	No	Parameters	Units	Source	
Cross-sectional area of the i <sup>th</sup>	$Ai = \pi \left( dx/2 \right)^2$	Eq. 1	<i>Ai</i> : cross-sectional area of the i <sup>th</sup> root at a given	mm <sup>2</sup>	Gonzalez-Ollauri	and
root at a given intersection			intersection plane		Mickovski (2016)	
plane			<i>dx</i> : root diameter	mm		
Root cross-sectional area at a	$Ar(z_i) = \Sigma Ai$	Eq.2	$Ar(z_i)$ : cross-sectional area of all roots crossing a	mm <sup>2</sup>	Gonzalez-Ollauri	and
given intersection plane			given intersection plane		Mickovski (2016)	
Vertical root distribution	$Ar(z) = Aro. e^{\frac{-z}{b}}$	Eq. 3	Ar(z): cross-sectional area of all roots along the	mm <sup>2</sup>	Preti et al. (2010)	
(VRD)			soil profile			
			Aro: cross-sectional area of the plant stem above	mm <sup>2</sup>		
			the root collar			
			<i>b</i> : mean rooting depth	mm		
			z: soil depth	mm		
Mean rooting depth in arid	$b = - \frac{\alpha}{2\alpha}$	Eq. 4	$\alpha$ : mean precipitation intensity per event over the	mm	Laio et al. (2006)	
and semi-arid ecosystems	$b = \frac{\alpha}{n(\theta f c - \theta w p)(1 - \frac{\lambda \alpha}{E t p})}$		growing season	event-1		
			n: soil porosity			
				ppu		
			capacity			
				ppu		

$\Delta r_p:$ columetric soil moisture content at withing point $\lambda:$ frequency of precipitation events over the growing season $Ep:$ total potential evapotranspiration over the growing seasonppu events mmMean rooting depth in humid ecosystems $b = \frac{\alpha}{n(\theta f c - \theta w p)}$ Eq. 5Image: Columetric soil moisture content at withing growing seasonmmMean rooting depth in humid ecosystems $b = \frac{\alpha}{n(\theta f c - \theta w p)}$ Eq. 5Image: Columetric soil moisture content at withing growing seasonmmMater available to plants in the soil $WAP = \theta f c - \theta w p$ Eq. 6 $WAP:$ water available to plantsppuArea at the root collar the soil $Aro = \frac{Mr}{b \rho r}$ Eq. 7 $Aro:$ cross-sectional area of the plant stem above the root collar $M:$ plant belowground biomassmn² g g g mm³Preti et al. (2010)	[]						
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Area at the root collar $Aro = \frac{Mr}{b \rho r}$ Eq. 7 $Aro:$ cross-sectional area of the plant stem above the root collar $Mr:$ plant belowground biomass $mm^2$ Preti et al. (2010)	Water available to plants in	$WAP = \theta fc - \theta wp$	Eq. 6	<i>WAP</i> : water available to plants	ppu		
$Aro = \frac{1}{b \rho r}$ the root collar $Mr: \text{ plant belowground biomass}$ g	the soil						
the root collar <i>Mr</i> : plant belowground biomass g	Area at the root collar	$Aro = \frac{Mr}{h cm}$	Eq. 7	Aro: cross-sectional area of the plant stem above	mm <sup>2</sup>	Preti et al. (2010)	
		ט ט		the root collar			
pr: root mass density g mm <sup>-3</sup>				Mr: plant belowground biomass	g		
				pr: root mass density	g mm <sup>-3</sup>		
Growing-degree day $GDDi = \frac{Tmx - Tmn}{2} - Tbase$ Eq.8 $Tmx$ : daily maximum air temperature °C McMaster and Wilhelm	Growing-degree day	$GDDi = \frac{Tmx - Tmn}{2} - Tbase$	Eq.8	<i>Tmx:</i> daily maximum air temperature	°C	McMaster and V	Vilhelm
<i>Tmn</i> : daily minimum air temperature (1997)		-		<i>Tmn</i> : daily minimum air temperature		(1997)	

	$\begin{cases} \sum_{i=1}^{n} GDDi \ge 200^{\circ} \text{C. VS start} \\ Tsoil \le 4^{\circ} \text{C VS end} \end{cases}$		<i>Tbase:</i> optimum daily mean temperature for plant growth <i>i</i> : i <sup>th</sup> day <i>Tsoil</i> : daily mean soil temperature	°C	
Aridity index	$AI = \frac{Etp}{PCP}$	Eq. 9	<i>PCP</i> : total precipitation over the growing season	mm	Greve et al. (2019)
Potential evapotranspiration	$Etp = 0.00128 \frac{Rnl}{58.3} \frac{\Delta}{\Delta + \gamma}$	Eq. 10	<i>Rnl</i> : net solar radiation Δ: slope of saturation vapour pressure γ: psychrometric constant	MJ m <sup>-2</sup> day <sup>-1</sup> kPa °C kPa °C	Priestly and Taylor (1972)
Soil moisture at field capacity	$\theta f c = \theta_{33} + 1.23 \theta_{33}^2 - 0.374 \theta_{33} - 0.015$ $\theta_{33} = -0.251Sa + 0.195Cl + 0.011SOC + 0.006Sa.SOC - 0.027Cl.SOC + 0.452Sa.Cl + 0.299$	Eq. 11	<ul> <li>θ33: soil moisture at -33 kPa of matric suction</li> <li>Sa: sand content in the soil</li> <li>Cl: clay content in the soil</li> <li>SOC: soil organic carbon</li> </ul>	ppu ppu ppu ppu	Toth et al. (2015)
Soil moisture at wilting point	$\theta_{wp} = \theta_{1500} + 0.14\theta_{1500} - 0.02$ $\theta_{1500} = -0.024Sa + 0.487Cl + 0.006SOC$ + 0.005Sa.SOC - 0.013Cl.SOC + 0.068Sa.Cl + 0.031	Eq. 12	$\theta$ 1500: soil moisture at -1500 kPa of matric suction	рри	Toth et al. (2015)





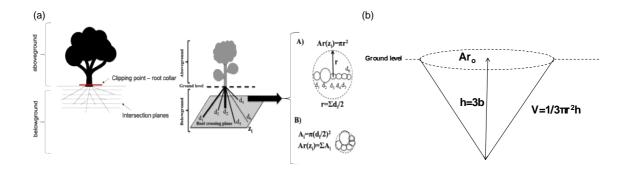




Figure 3. (a) Illustration of the methodological approach followed to separate aboveground and belowground plant parts and to describe vertical root distribution (VRD) for the studied plant species. Approach B was followed to describe the crosssectional area of all roots at a given crossing plane (Ar(z)), as it does not overestimate the root cross-sectional area compared to approach A. r stands for the rooting radius and d for the diameter of ith root at a given crossing plane. (b) VRD was modelled as a cone with base's area Aro (i.e. root collar area; mm<sup>2</sup>) and height 3b, being b the mean rooting depth (mm) and 3b the soil depth at which 95% of the roots are found (Laio et al., 2006). V stands for the cone volume and h for the cone height. See mathematical formulation in Supplementary Material.

295 2.5. Relationship between soil and plant attributes with key VRD parameters

296

297 We investigated the relationship between the studied soil (Section 2.3) and plant attributes (Section 2.4) with the relevant/key parameters used to portray VRD (Aro: cross-sectional area 298 299 of the root collar, and b: mean rooting depth) through a data mining workflow (Supplementary 300 Material – Fig. S1) which was built using the statistical language R v5.5.1 (R Core team, 2018). 301 We also included plant belowground biomass (Mr) in the analysis, as Mr will ultimately limit 302 the extent of VRD (Gonzalez-Ollauri and Mickovski, 2016). This workflow was used to 303 accomplish three objectives: (i) to evaluate the ability to predict relevant VRD parameters using the investigated soil and plant attributes as predictors, (ii) to evaluate the importance of 304

ach plant and soil predictor on the relevant VRD parameters, and (iii) to evaluate predictorresponse dependency - i.e. how the response variable changes following predictor changes.

308 To accomplish objective (i), 100 random forest models (RF; Breiman, 2001) were fitted with 309 1000 regression trees each, using the R package "randomForest" (Liaw and Wiener, 2020). 310 Only uncorrelated attributes to the VRD parameters were considered to fit RF models. Each 311 RF model was cross-validated with a bootstrapping method without replacement (e.g. 312 Gonzalez-Ollauri et al., 2020) and through the evaluation of the coefficient of determination (R<sup>2</sup>), root mean square error (RMSE) and variance explained (VExp) following the least-313 314 squares method (Fuller, 1987). The pool of cross-validation coefficients retrieved from 315 implementing the data mining workflow was examined by plotting their corresponding 316 probability density functions (Appendix A).

317

To accomplish objective (ii), the importance of each plant and soil attribute on the response variables (i.e. key VRD parameters) was examined on the basis of permutation tests using the R package "caret" (Khun et al., 2018), which measures attribute importance by observing model performance when each predictor is randomly dropped out from fitting a RF model during the training step (e.g. Strobl et al., 2008).

323

To achieve objective (iii), we examined the Partial Dependence Plots (PDPs; Hastie et al., 2009) retrieved from using the R package "pdp" (Greenwell, 2017). PDPs were retrieved to show whether the interaction between a target VRD parameter and a target plant and soil attribute was linear, monotonic, or more complex in the fitted RF models, representing how a given attribute influenced the prediction on average for a given VRD parameter. The R script used to implement the data mining workflow described above is provided in Appendix C.

- 331 2.6. Empirical vs. parametric, ecohydrological model for vertical root distribution
- 332

333 We tested the predictive capacity of an existing parametric ecohydrological model for vertical 334 root distribution (Eqs. 4-5, Table 1; Laio et al. 2006; Preti et al., 2010; Gonzalez-Ollauri and 335 Mickovski, 2016) against the empirical VRD model fitted to the measured data described in 336 Section 2.4. The parametric ecohydrological VRD model was firstly developed for arid and 337 semi-arid ecosystems by Laio et al. (2006) and extended by Preti et al. (2010), and then adapted 338 to temperate-humid climates by Gonzalez-Ollauri and Mickovski (2016). This model estimates 339 the mean rooting depth (b; Eqs. 4 and 5; Table 1) on the basis of pedo-climatic parameters (i.e. 340  $\alpha$ : mean precipitation depth during the growing season;  $\lambda$ : frequency of precipitation events; 341 *Etp*: potential evapotranspiration; Section 2.2) and of the product between soil porosity (n) and 342 the water available to plants in the soil (WAP; Eq. 6; Table 1), function of the difference 343 between the volumetric soil moisture content at field capacity ( $\theta_{fc}$ ) and at wilting point ( $\theta_{wp}$ ). 344 Different equations for b must be considered depending on whether the aridity index (AI; Eq. 345 8) is greater than 1 (i.e. arid climate; Eq. 4; Preti et al., 2010) or lower than 1 (i.e. humid 346 climate; Eq.5; Gonzalez-Ollauri and Mickovski, 2016). VRD is then modelled with a negative 347 exponential equation (Eq. 5; Table 1) using b, the cross-sectional area at the root collar (Aro) 348 and the soil depth (z; mm) as inputs, assuming that the probability density function for the daily 349 rainfall intensity at the study site is exponentially distributed (Laio et al., 2006). The cross-350 sectional area at the root collar (Aro) is estimated using plant-specific information and the 351 rooting depth under the assumption that the distribution of root biomass along the soil profile 352 can be portrayed with a conical-shape-volume (Fig. 3b; Section 2.4; Eq. 6; Table 1; 353 supplementary material). In addition, we assumed that the portion of soil explored by roots was 354 uniform and isotropic.

356 The outcomes from the parametric, ecohydrological and the empirical VRD models were 357 compared in the light of the outputs for the root cross-sectional area (Ar) of all roots along the 358 soil profile. To do so, Ar was firstly retrieved for soil depths of 0 mm - 100 mm using the parametric ecohydrological and empirical VRD models (Section 2.4), respectively. Then, the 359 360 two Ar datasets were log-transformed and plotted together to graphically evaluate the 361 mathematical relationship between the two models. Subsequently, a linear regression model 362 was fitted between the two retrieved, log-transformed Ar datasets in R v3.5.1. Additionally, 363 the correlation between the linear fitting parameters and the studied plant and soil attributes 364 was examined by estimating pairwise Pearson's correlation coefficients.

365

366	2.7.	Statistical	analysis

367

368 Normality checks were undertaken for every studied soil and plant attribute with the Shapiro 369 Wilk test. Soil and plant attributes were aggregated into plant species and study site, 370 respectively, for statistical analysis. Statistical differences in plant attributes between plant species and investigation site were evaluated with the non-parametric Kruskal Wallis ( $\chi^2$ ) test, 371 as plant attributes did not follow a normal distribution. Where statistically significant 372 373 differences were encountered, plant attribute's differences between two plant species were 374 evaluated with the non-parametric Wilcoxon (W) test. Statistical differences in soil attributes 375 between plant species and investigation site were evaluated with one-way ANOVA (F) and Kruskal Wallis ( $\chi^2$ ) tests for normal and non-normal distributed variables, respectively. 376 Vertical root distribution (VRD) differences between investigated sites were evaluated per 377 plant species with the Kruskal Wallis ( $\chi^2$ ) test, as VRD did not follow a normal distribution. 378 379 Where statistically significant differences were encountered, the differences within the plant

380 species were evaluated with the non-parametric Wilcoxon (W) test. VRD differences between 381 plant species for a given investigation site were examined with the same approach indicated 382 before. Statistical differences between the importance of the attributes used as predictors for 383 the selected VRD parameters were also evaluated with the Kruskal Wallis test. The statistical 384 relationship between the studied plant and soil attributes was evaluated with the pair-wise 385 Pearson's correlation test and interpreted on the basis of the resulting correlogram plot. All statistical tests were carried out at the 95 % and 99 % confidence level using the 'base' package 386 387 embeddedw in the statistical computing software R v3.5.1 (R Core Team, 2018).

388

389 3. Results

390

- 391 3.1. Climate attributes
- 392

The growing season duration for the period 2005-2019 was on average 50±18 days long, generally starting in early July and ending in early September. The aridity index of the study site during the snow-free period was 0.91±0.2, indicating that the temperate-humid, ecohydrological VRD model (Eq.5; Table 1) must be implemented for the study area. The mean precipitation depth per event during the growing season ( $\alpha$ ) was 6.68±2.33 mm, the frequency of precipitation events ( $\lambda$ ) was 0.56±0.05, and the total potential evapotranspiration (*Etp*) during the vegetative season was 68.8±2.1 mm.

400

401 3.2.Soil attributes

402

403 The examined soil attributes (Table 2) differed significantly between the investigated study 404 sites. Site 3 had a substantially higher slope gradient ( $\chi^2$ :18.2 df:2 p<0.01), it was consistently

- 405 concave in terms of the terrain curvature, and it had a more South-facing aspect than Sites 1 406 and 2. In addition, Site 3 had a significantly higher soil moisture (F:65.89 df:1 p<0.01), soil 407 organic carbon (F:45.3 df:1 p<0.01), and proportion of fine soil materials ( $\chi^2$ :35 df:2 p<0.01)
- 408 than Sites 1 and 2, while having a substantially lower bulk density ( $\chi^2$ :20.2 df:2 p<0.01), soil
- 409 skeleton ( $\chi^2$ :35 df:2 p<0.01), and coarse soil materials ( $\chi^2$ :35 df:2 p<0.01) than the other two

Table 2. Soil attributes investigated in this study averaged per plant species and per sampling site ± standard deviation. θg: gravimetric moisture content; ρ<sub>bk</sub>: dry bulk density; n: soil porosity;
 SOC: soil organic carbon; Sk: soil skeleton.

		Slope (°)	Curvature	Aspect	<i>θg</i> (%)	$ \rho_{bk} (\text{g cm}^{-3}) $	n	<i>SOC</i> (%)	рН	Sk (%)
Plant species	Euphrasia									
	minima	8.42±8.7	convex	145.42±22.8	29.20±11.4	1.18±0.2	0.56±0.1	7.70±2.6	4.50±0.1	20.70±7.2
	Leucanthemopsis									
	alpina	8.42±6.3	flat/convex	145±25.3	27.46±9.3	1.24±0.2	0.53±0.1	6.59±1.4	4.50±0.1	20.70±7.2
	Poa alpina	11.50±7.3	flat/convex	152.08±18.8	27.43±9.2	1.20±0.2	0.55±0.1	7.83±2.6	4.50±0.1	20.70±7.2
Sampling site	Site 1	6.92±4.5	flat/convex	133.50±11.1	19.44±7.1	1.37±0.2	0.48±0.1	5.52±1.4	4.60±0.0	16.30±0.0
	Site 2	4.33±2.1	convex	135.5±9.81	26.05±4.0	1.26±0.1	0.52±0.0	6.96±1.4	4.40±0.0	30.40±0.0
	Site 3	17.08±7.3	concave	173.5±14.9	38.60±5.5	0.98±0.2	0.63±0.1	9.64±1.6	4.50±0.0	15.40±0.0

416 Table 2 Cont. Soil attributes investigated in this study averaged per plant species and per sampling site ± standard deviation. θfc: volumetric soil moisture content at field capacity; θwp:

#### 417 volumetric soil moisture content at wilting point.

		Clay (%)	Fine silt (%)	Coarse silt	Fine sand	Coarse sand	<i>θfc</i> (%)	<i>θwp</i> (%)
				(%)	(%)	(%)		
Plant species	Euphrasia							
	minima	1.40±0.34	5.67±1.11	11.30±3.33	50.73±3.64	30.87±2.15	27.851±0.3	1.56±0.04
	Leucanthemopsis							
	alpina	1.40±0.34	5.67±1.11	11.30±3.33	50.73±3.64	30.87±2.15	28.33±0.2	2.01±0.10
	Poa alpina	1.40±0.34	5.67±1.11	11.30±3.33	50.73±3.64	30.87±2.15	29.18±0.4	3.09±0.20
Sampling sites	Site 1	1.00±0.0	4.30±0.0	9.20±0.0	52.30±0.0	33.20±0.0	28.78±0.7	2.37±0.9
	Site 2	1.40±0.0	5.80±0.0	15.80±0.0	45.90±0.0	31.20±0.0	28.10±0.6	2.20±0.7
	Site 3	1.80±0.0	6.90±0.0	8.90±0.0	54.00±0.0	28.20±0.0	28.23±0.9	2.16±0.7

419	investigated sites (Table 2). However, significant differences for the investigated soil attributes
420	were not detected between the three studied plant species (Table 2; F:1.1 df:2 p=0.34). The
421	available water to plants in the soil (Eq. 6, Table 1) was on average of 26.13±0.3 %.
422	

3.3.Plant attributes

424

425 The evaluated plant attributes (Table 3) were statistically different between the three studied 426 plant species. In particular, Poa alpina individuals had a substantially larger crown spread (CS;  $\chi^2$ :26.7 df:2 p<0.01), plant projected area (*Sp*;  $\chi^2$ :26.7 df:2 p<0.01), cross-sectional area at the 427 root collar (Aro;  $\chi^2$ :27.1 df:2 p<0.01), aboveground biomass (Ma;  $\chi^2$ :26.4 df:2 p<0.01), and 428 429 root biomass (*Mr*;  $\chi^2$ :29.3 df:2 p<0.01) than those of the other two studied species (Table 3). 430 However, the measured mean rooting depth (b) was not statistically different between the three studied plant species (Table 3;  $\chi^2$ :0.6 df:2 p=0.76). On other hand, the three investigated study 431 sites did not present statistical differences in terms of the evaluated plant attributes ( $\chi^2$ :0.3 df:2 432 p=0.85) with the exception of the observed b, which was significantly higher in Site 3 than in 433 Sites 1 and 2 (Table 3;  $\chi^2$ :26.7 df:2 p<0.01). 434

435

436 3.4. Vertical root distribution

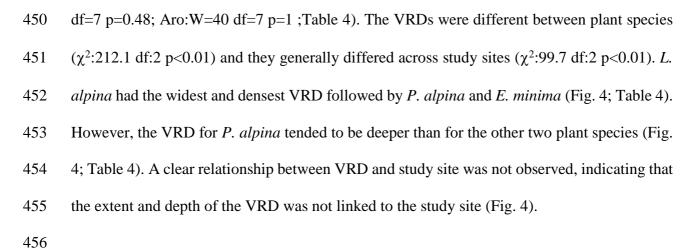
437

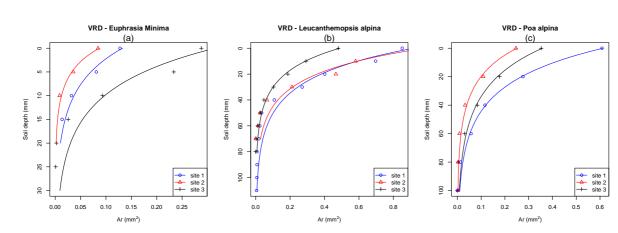
The measured vertical root distribution (VRD) decreased exponentially with soil depth for the three studied plant species and at the three study sites (Fig 5). As a result, negative exponential models (Eq. 3. Table 1) were successfully fitted to the observed data (i.e. empirical model) with high goodness of fit ( $R^2$ >0.9) in all cases. The model fitting parameters (Table 4) did not differ statistically from the measured VRD parameters (b:W=49 443 Table 3. Plant attributes investigated in this study averaged per plant species and per sampling site ± standard deviation . CS: crown spread; Sp: plant's aerial projected area; Aro: root collar 444 area; b: mean rooting depth; Ma: aboveground biomass; Mr: root biomass; ALR: allometry ratio -i.e. Mr/Ma.

		CS (cm)	Sp (cm <sup>2</sup> )	Aro (mm <sup>2</sup> )	<i>b</i> (mm)	Ma (mg)	<i>M</i> r (mg)	ALR
Plant species	Euphrasia minima	1.60±1.4	3.44±7.9	0.17±0.1	15.00±10.9	13.38±12.1	1.08±1.1	0.12±0.2
	Leucanthemopsis alpina	3.84±0.8	12.09±4.9	0.38±0.1	14.44±10.9	85.43±40.8	62.12±21.0	0.82±0.4
	Poa alpina	8.01±3.0	56.90±45.7	1.15±0.7	15.83±8.1	226.42±162.4	379.07±366.3	1.82±1.2
Sampling sites	Site 1	4.06±3.3	20.99±31.4	0.72±0.9	4.72±1.9	100.10±132.3	189.71±371.1	1.09±1.1
	Site 2	3.90±2.2	15.49±13.8	0.45±0.4	16.67±7.8	133.03±173.3	158.17±274.4	0.95±1.1
	Site 3	5.49±4.1	35.94±50.5	0.52±0.5	23.89±6.0	92.11±72.5	94.39±142.9	0.71±0.9

446<br/>447<br/>448Table 24. Vertical root distribution (VRD) parameters measured, retrieved though fitting nls exponential models to the measured data -i.e. empirical VRD model (†), and predicted with the<br/>parametric, ecohydrological VRD model (†) per plant species and sampling site. b: mean rooting depth; Aro: cross-sectional area at the root collar.448

		Site 1					Site 2						Site 3					
Plant	<i>b</i> (mm)	$b^{\dagger}$	$b^{\dagger \ \dagger}$	Aro	Aro <sup>†</sup>	$Aro^{\dagger\dagger}$	<i>b</i> (mm)	$b^{\dagger}$	$b^{\dagger \ \dagger}$	Aro	$Aro^{\dagger}$	$Aro^{\dagger \dagger}$	<i>b</i> (mm)	$b^{\dagger}$	$b^{\dagger\dagger}$	Aro	$Aro^{\dagger}$	$Aro^{\dagger\dagger}$
species		(mm)	(mm)	(mm <sup>2</sup> )	(mm <sup>2</sup> )	(mm <sup>2</sup> )		(mm)	(mm)	(mm <sup>2</sup> )	(mm <sup>2</sup> )	(mm <sup>2</sup> )		(mm)	(mm)	(mm <sup>2</sup> )	(mm <sup>2</sup> )	(mm <sup>2</sup> )
Е.	4.17±0.9	7.89	52.7	0.13±0.0	0.13	0.013	17.50±12.9	5.20	49.9	0.08±0.0	0.08	0.016	23.3±3.8	8.50	38.1	0.29±0.1	0.30	0.09
minima																		
L.	3.33±0.0	22.80	56.9	0.38±0.1	0.9	1.89	15.00±4.3	19.70	49.9	0.38±0.1	1.00	2.24	25.0±10	19.10	42.8	0.37±0.1	0.50	1.66
alpina																		
Р.	6.67±1.9	24.4	2.6	1.66±1.0	0.60	15.88	17.50±5.7	21.70	47.9	0.88±0.2	0.20	12.68	23.33±3.8	26.60	42.5	0.91±0.6	0.40	8.33
alpina																		





458

Figure <u>34</u>. Vertical root distribution (VRD) for (a) Euphrasia minima, (b) Lecanthemopsis alpina, and (c) Poa alpina.
Triangles, dots and crosses represent observed values for the root cross-sectional area of all roots found at a given soil depth as described in Section 2.4. The lines portray the nls exponential models fitted to the measured data points -i.e.
empirical VRD model. See online version for colours.

463

464

465

The cross-validation results from fitting random forest models to the key vertical root distribution (VRD) parameters (i.e. *b*, *Aro*, and *Mr*; Appendix A - Table A1 and Figure A1; Supplementary Material) suggested that the latter can be predicted successfully using the studied plant and soil attributes as predictors.

<sup>3.5.</sup>Influence of plant and soil attributes on key vertical root distribution parameters

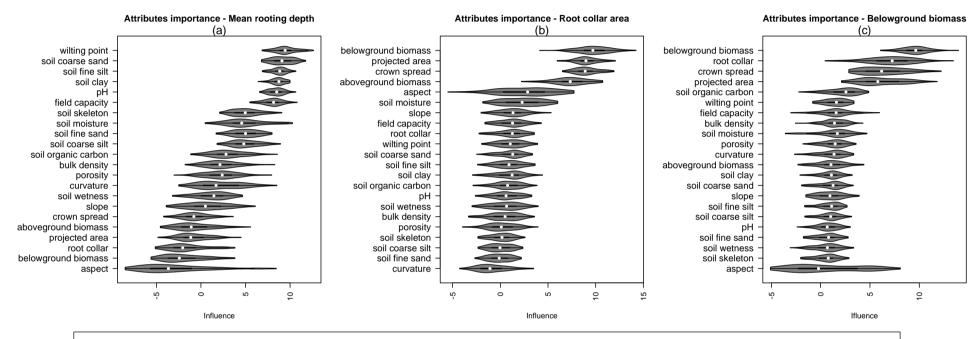


Figure 45. Violin plots depicting the influence of the studied plant and soil attributes on relevant vertical root distribution (VRD) parameters -i.e. (a) mean rooting depth, b; (b) root collar area, Aro; (c) plant belowground biomass, Mr. The white dot within the violin plot boxes represent the median while the grey area around the box represents the probability density of the data at different values.

472 The mean rooting depth (b) was chiefly influenced by soil attributes (Fig. 5a) such as the soil 473 volumetric moisture content at wilting point, followed by the soil texture (Fig. 5a). Soil pH and 474 the soil moisture at field capacity also appeared to significantly influence b (Fig. 5a). In the 475 light of the correlogram (Fig. 6), the rooting depth was strongly correlated to the soil texture, 476 positively with clay and silt contents and negatively with coarse sand content. The slope aspect, 477 the soil moisture, and organic carbon content also had a strong, positive correlation with b (Fig. 478 6). In the light of the PDPs (Appendix B - Fig. B1), there was a positive influence of the wilting 479 point on b (i.e. the higher  $\theta wp$  the higher b), up to 3.4 %, after which a constant effect was 480 observed. Field capacity also had a positive effect on b only detected when  $\theta fc$  was above 15.6 481 %. We noticed a negative influence of soil pH on b when the former was above 4.5. The 482 attributes that did not have a substantial effect on b (Fig. 5a) had a remarkable effect when the 483 PDPs were assessed (Appendix B - Fig. B1). For example, deeper root systems were found 484 under steeper conditions. However, shallower root systems were encountered when soil was 485 wetter, but shifts in the soil hydrological regime led to changes in the influence of soil moisture 486 on b. A negative effect of soil moisture on b was observed under residual and saturated regimes, 487 and a positive effect was noticed under the transitional regime. Contrariwise, soil porosity and 488 organic carbon had a positive effect on b (Tables 2 and 3; Fig. 6) which was not detected in the 489 PDPs for the soil porosity (Fig. B1).

490

The cross-sectional area at the root collar (*Aro*) was mostly affected by plant attributes (Fig. 5b). The root biomass, surface projected area, crown spread, and aboveground biomass had a significant influence on *Aro*. However, the investigated soil attributes did not have a substantial influence on *Aro* on the basis of the RF model outputs (Fig. 5b). In the light of the correlogram plot (Fig. 6), *Aro* had a strong, positive correlation with above- and belowground biomass but also with the soil water content at field capacity and wilting point. According to the PDPs (Appendix B - Fig. B2), all the examined plant attributes had a consistent, positive effect on *Aro*. We also observed that some soil attributes had a consistent effect on *Aro* when the PDPs were examined (Fig. B2) that could not be detected in the correlogram (Fig. 6) or in relative influence plots (Fig. 5b). According to the PDPs, soil porosity, the percentage of clay, fine silt and fine sand had a negative effect on *Aro*, while the percentage of coarse sand and coarse silt had a positive effect. We also observed that *Aro* tended to be wider under steeper slope conditions, and narrower when soil organic carbon increased (Fig. B2).

504

505 The root biomass (Mr) was predominantly influenced by plant attributes (Fig. 5c) such as the 506 aboveground biomass, Aro, crown spread, and plant's surface projected area. However, the 507 influence of soil organic carbon on Mr was significantly higher than the influence of the rest 508 of the studied soil attributes (Fig. 5c). In the light of the correlogram plot (Fig. 6), Mr had a 509 strong, positive correlation with the aboveground biomass, the crown spread, and Aro. In 510 addition, soil attributes, such as the soil moisture at field capacity and wilting point, also had a 511 strong positive correlation, with the latter being the attribute with highest correlation to Mr 512 (Fig. 6). According to the PDPs (Appendix B - Fig. B3), all the plant attributes studied had a consistent, positive effect on Mr. In addition, we noticed a consistent effect of some of the soil 513 514 attributes studied on Mr which resembled the effects observed for Aro (Fig. B2).

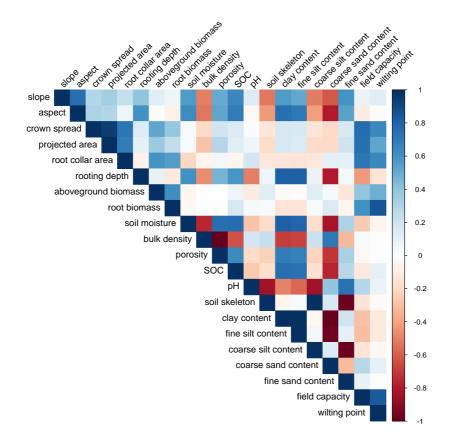


Figure <u>56</u>. Correlation plot depicting Pearson's correlation coefficient between the investigated plant and soil attributes.
 Blue colour: positive correlation; Red colour: negative correlation. The darker the colour shade, the higher the correlation
 between two attributes. See online version for colours.

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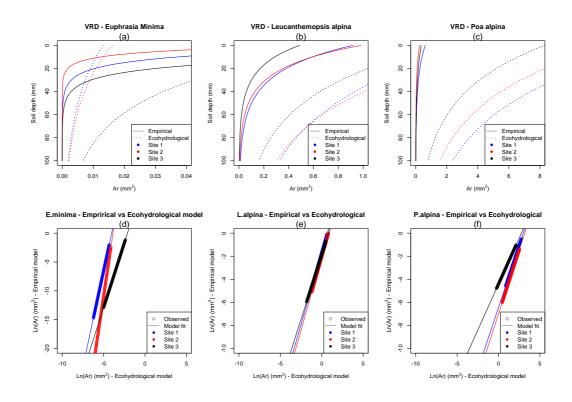
#### 521 3.6.Ecohydrological model for vertical root distribution

522

We found a substantial mismatch between the chosen parametric ecohydrological and the 523 empirical VRD models (Figs. 8a-c; Table 4). This showed that the existing parametric, 524 ecohydrological VRD model cannot readily predict VRD under the pedoclimatic conditions of 525 the study area. However, we detected a consistent ( $R^2 > 0.9$ ) linear relationship between the 526 527 ecohydrological and empirical VRD models when Ar(z) was log-transformed (Figs. 8d-e) -i.e. 528 a consistent exponential fit was observed when comparing empirical, untransformed Ar(z)529 values against parametric, untransformed Ar(z) values. The fitting parameters for the log-530 transformed linear models established between ecohydrological and empirical VRD models

are shown in Table 5. We observed a strong correlation (r > 0.5) between most of the studied plant attributes and the fitting parameters for the log-transformed linear models (Table 6). In addition, a strong correlation was observed between the fitting parameters and the soil moisture content at field capacity and at wilting point (Table 6).

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- 536



538Figure 67. (a-c) Vertical root distribution (VRD) models fitted with a nls exponential model to the measured data – i.e.539empirical VRD model (solid lines) and with the parametric, ecohydrological VRD model (dotted line) (d-e) Mathematical540relationship between empirical and ecohydrological VRD models established with a log-transformed linear model of the541form "Ln(Empirical Ar(z))=A+BxLn(Ecohydrological Ar(z))" (see Table 5 for fitting parameters). Also, see online version of542this manuscript for colours.

### 543<br/>544Table 35. Fitting parameters for the log-transformed linear model the form "Ln(Empirical Ar(z)) = A + BxLn(Ecohydrological Ar(z))"544Ar(z)" fitted between empirical and ecohydrological VRD models.

	Si	te 1	Sit	e 2	Site 3		
Plant species	А	В	А	В	А	В	
E. minima	26.88	6.67	37.46	9.68	9.47	4.47	
L. alpina	-1.67	2.49	-2.07	2.53	-1.86	2.24	
P. alpina	-6.44	2.15	-7.00	2.21	-4.41	1.59	

 Table <u>46</u>. Pearson's correlation coefficients for the pair-wise correlation between the evaluated soil and plant attributes and the fitting parameters for the log-transformed linear model the form "Ln(Empirical Ar(z))=A+BxLn(Eco-hydrological Ar(z))" fitted between empirical and ecohydrological VRD models.

	1 .	1
Variable	А	В
Slope	-0.36	-0.42
Aspect	-0.31	-0.36
Curvature	-0.05	0.03
CS	-0.74	-0.73
Sp	-0.56	-0.58
Aro	-0.65	-0.60
Ма	-0.75	-0.68
Mr	-0.61	-0.54
b	-0.11	-0.11
θg	0.20	0.23
ры	-0.12	-0.14
n	0.18	0.20
SOC	-0.18	-0.20
рН	-0.12	-0.14
sk	-0.09	-0.16
Clay	0.19	0.29
Fine silt	-0.14	-0.16
Coarse silt	-0.12	-0.13
Fine sand	0.19	0.29
Coarse sand	-0.21	-0.31
θfc	0.16	0.19
θwp	-0.59	-0.57
L	I	l

#### 550 4. Discussion

551 4.2. Vertical root distribution

552 The vertical root distribution (VRD) of the three studied alpine plant species decreased 553 exponentially with soil depth. Accordingly, VRD was successfully described with a negative 554 exponential model that was fitted to the measured data (Fig. 4). This is consistent with VRDs 555 reported for shrub, woody, and herbaceous plant species in Mediterranean (Preti et al., 2010; 556 Tron et al., 2014), southern alpine (Burylo et al., 2011) and temperate-humid ecosystems 557 (Gonzalez-Ollauri and Mickovski, 2016; Tardio et al., 2016), and it confirms that the proposed 558 approach for describing VRD in herbaceous plants is methodologically robust across terrestrial 559 ecosystems in Europe.

560

561 We attributed the observed VRD differences across plants and sites (Fig. 4) to the differences in plant attributes that we found (Table 3). In spite of the differences observed in key VRD 562 563 parameters across sites (Table 3), and of the strong influence of site-specific attributes on the 564 key VRD parameters that we noticed herein (Fig. 5 and Section 4.2), the direction of the effect 565 of the study site on the size and depth of the measured VRDs was unclear, given that the extent 566 of VRD changed with the plant species from site to site (Fig. 4). Yet, there were limitations to 567 the study, as we did not evaluate altitudinal differences between plant individuals (e.g. Gale, 568 2004; Miyamoto et al., 2015), nor the differences in soil nutrients between sampling locations 569 (e.g. Forde and Lorenzo, 2001), or the climatic differences across study sites (Schenk and 570 Jackson, 2005). We think that all these aspects deserve further consideration to expand on our 571 findings related to how VRD is shaped by local site conditions.

573 The proposed VRD description approach has been used effectively in woody plant species 574 under both sloped and flat topographies (Tron et al., 2014; Tardio et al., 2016; Gonzalez-Ollauri 575 et al., 2020). Still, its ability to capture realistically large roots (> 3 mm in diameter, incl. tap 576 roots) in woody plants needs further verification, notwithstanding the fact that the *in situ* 577 description of root systems for woody plants is methodologically challenging (Böhm, 1979), 578 and it generally focuses on one or two vertical profiles of the root system (e.g. Tardio et al., 579 2016; Gonzalez-Ollauri et al., 2020) from which it is hard to comprehensively capture root 580 system features. However, the VRD description approach followed herein is methodologically 581 simple and easy to implement, it provided a good and realistic picture of the VRD for the 582 studied alpine plants with root systems mainly comprising fine roots, and it generated 583 information directly applicable in workflows needing VRD information such as plant-water 584 uptake models (e.g. Laio, 2006; Shukla, 2014) or soil-root reinforcement estimation 585 approaches (e.g. Gonzalez-Ollauri and Mickovski, 2016, 2017b; Kokutse et al., 2016).

586

4.3.Influence of climate, plant and soil attributes on key vertical root distribution parameters

The rooting depth (*b*) and root biomass (*Mr*) for the three studied plant species (Table 3) were within the ranges described in Pohl et al. (2011) and Hudek et al. (2017) for alpine ecosystems. However, these were below the ranges reported for semi-arid (Preti et al., 2010) and temperatehumid (Gonzalez-Ollauri and Mickovski, 2016) climates. These differences were attributed to the short duration of the growing season in our study area, and associated to long periods with snow cover and low temperatures, which likely limited plant development (e.g. Kaspar and Bland, 1992; Lahti et al., 2005; Alvarez-Uria and Körner, 2007).

The key studied VRD parameters (i.e. *b*: mean rooting depth; *Aro*: cross-sectional area at the root collar; and *Mr*: root biomass) were distinctly influenced by the investigated soil and plant attributes (Figs 6, 7 and Appendix B – Figs. B1-B3). We observed that, while the rooting depth was mostly site-specific, the allocation of biomass to the belowground plant parts and its distribution along the soil profile was both species-specific and reliant on relevant soil ecohydrological features.

- 603
- 604 4.3.1. Rooting depth
- 605

606 The mean rooting depth (b; Laio et al., 2006) was chiefly influenced by soil attributes 607 governing the water available to plants in the soil (WAP; i.e. difference between soil water 608 content at field capacity ( $\theta fc$ ) and wilting point ( $\theta wp$ ); Fig. 5a, 7 and Appendix B – Fig. B1; 609 Table 4; Eq.4 Table 1; Casper et al., 2003). It is worth noting that  $\theta fc$  and  $\theta wp$  were estimated 610 in the light of well-established pedotransfer functions nested in the VRD model (Eqs. 4, 5, 11 611 and 12; Table 1) with relatively low sensitivity (Gonzalez-Ollauri and Mickovski, 2016) and 612 which employed soil texture, soil organic carbon, and porosity as inputs (Toth et al., 2015). 613 Accordingly, soils with high organic carbon and with a fine texture (i.e. high clay and silt 614 content) would have high water retention capacity (Kirkham, 2005; Lu and Likos, 2004) and, 615 as a result, deeper root systems, as it was shown herein (Figs. 6 and B1; Table 3; Schenk and 616 Jackson, 2005; Gonzalez-Ollauri and Mickovski, 2016). In fact, we observed clear differences 617 across sites in terms of the soil attributes governing WAP (Table 2), which may explain why 618 Site 3 had substantially deeper rooting depths than sites 1 and 2 (Table 3). The effect of the 619 soil's ecohydrological characteristics on the rooting depth has been highlighted in previous 620 studies (e.g. Schenk, 2005; Laio et al., 2006; Preti et al., 2010), suggesting that the ability of 621 roots to explore the soil in depth largely depends on the water mass balance within the topsoil

622 (Tsutsumi et al., 2003; Laio, 2006; Laio et al., 2006). In fact, the soil water mass balance 623 features in the proposed ecohydrological models predicting the mean rooting depth (Eqs. 4 and 624 5; Table 1; Laio et al., 2006; Gonzalez-Ollauri and Mickovski, 2016), and our results validate 625 that these models are conceptually correct. However, other models are nested into the 626 investigated VRD model (e.g. growing season duration, evapotranspiration, soil pedotransfer 627 functions, etc.), leading to likely propagation of errors and uncertainty (Taylor, 1997) that should be thoroughly investigated and dealt with prior to verifying the quality and robustness 628 629 of the VRD models (e.g. Gonzalez-Ollauri and Mickovski, 2017b).

630

631 Model differences between the options for arid or semi-arid (Eq. 4; Table 1) and temperate 632 humid (Eq.5; Table 1) climates imply rooting depth differences that were not tested herein. In 633 arid climates, water withdrawal through evapotranspiration limits the amount of water 634 available to plants in the topsoil, thus encouraging deeper rooting depths than in humid 635 climates. By contrast, in temperate humid climates, water inputs exceed outputs in the topsoil 636 (i.e. rainfall > evapotranspiration), leading to shallower rooting depths as roots do not face 637 water limitations in the topsoil - i.e. roots do not need to explore the soil in search of deep water 638 (Schenk and Jackson, 2005). In alpine climates, however, where snowfall, ground frost, and 639 short snow-free periods govern the soil ecohydrological behaviour (e.g. Molotch et al., 2009), 640 further model tuning is needed to fully capture the rooting depth and VRD with the proposed 641 parametric ecohydrological model (Fig. 7a-c; Table 4). In addition, the direct quantification of 642 soil attributes governing the available water to plants through, for example, retrieving the soil-643 water retention function (e.g. Zhang et al., 2019) or by evaluating the soil structure and 644 aggregates (e.g. Bengough, 2003) could shed more light on the effect of the soil's 645 ecohydrological characteristics on the rooting depth and VRD.

647 It is also worth noting the high influence of soil pH on the rooting depth (Fig. 4a; Fig. 6), which 648 on the basis of the correlogram (Fig. 6) and PDPs (Annex B - Fig.B1) was negative - i.e. 649 shallower rooting depths were noticed when the pH was higher. This observation may indicate 650 that plants tend to reduce root elongation and increase thickness as a strategy to promote 651 nutrient uptake when soil pH is high (e.g. Robles-Aguilar et al., 2019). The latter is somehow 652 supported by the positive interaction that we observed between soil pH and the cross-sectional area at the root collar (Aro; Figs. 6 and B2), suggesting that root thickness increased at the 653 654 collar when soil pH increased. Yet, as the variability in soil pH was small across study sites 655 (Table 3), we cannot convincingly elucidate possible reasons behind our observations and we 656 thus recommend further research on the effect of soil pH on the key VRD parameters.

657

658 In this study, there were also limitations to the analysis because site-specific records for the 659 climate attributes and for the snowpack depth, although capable of affecting the rooting depth in alpine ecosystems (e.g. Cooper et al., 2011), were not available. Additionally, we did not 660 661 consider the topographical effect of the slope gradient on the rainfall lost to runoff and, in turn, on the rooting depth (Tron et al., 2014; Tardio et al., 2016); nor the influence of preferential 662 flow paths in the soil (e.g. Clothier et al., 2008; Gonzalez-Ollauri et al., 2020) or the effect of 663 664 soil anisotropy on the rooting depth and VRD. We believe that all these aspects deserve detailed 665 consideration to improve the predictive capacity of the parametric VRD models studied here. 666 However, the consistent linear relationship between empirical and parametric ecohydrological VRD models reported in this study (Fig. 7d-e), and the high correlation found between fitting 667 668 parameters and soil attributes governing the available water to plants (i.e.  $\theta fc$  and  $\theta wp$ ; Table 669 6) set the direction of future research.

670

671 4.3.2. Cross-sectional area at the root collar and root biomass

673 The cross-sectional area at the root collar (Aro) and the root biomass (Mr) were mostly 674 influenced by the investigated plant attributes (Fig. 5 and Appendix B - Figs. B2 and B3). 675 However, a substantial effect of the soil water at field capacity and wilting point on Aro and 676 Mr was also detected (Fig. 6). It is worth noting that the combined effect of the soil and plant 677 attributes on the key VRD parameters was only evident when multiple data analysis approaches 678 were used together - i.e. relative influence from RF models (Fig. 5), pairwise correlation (Fig. 679 6) and PDPs (Appendix B), indicating that comprehensive data mining is needed to fully grasp 680 complex interactions between environmental variables affecting VRD (Supplementary 681 Material - Fig. S1). The findings shown herein corroborate that while the extent of the root system in the soil (i.e. rooting depth; Section 4.2.1) is delimited by the soil water mass balance 682 683 and its contributing soil attributes (i.e. soil texture, SOC,  $\theta fc$  and  $\theta wp$ ; Figs. 5a and 6), the plant 684 biomass allocated belowground is distributed along the VRD profile in the light of both plant-685 specific and soil attributes (Figs. 6b-c and 7). The latter aspect features in the proposed VRD 686 model through Aro (Eq.3; Table 1), which acts as scaling factor in the distribution of root 687 biomass along the root profile in the soil (Preti et al., 2010; Gonzalez-Ollauri and Mickovski, 688 2016). In this regard, we noticed that the magnitude of Aro was positively influenced by the 689 slope gradient whilst being negatively affected by SOC (Fig. B2). A plausible explanation for 690 the former is that plants tend to adopt anchoring strategies in steep slopes (e.g. Tardio et al., 691 2016), which may imply the allocation of root biomass near the ground surface to promote 692 anchorage to the ground and plant stability (Chiatante et al., 2003). In fact, we also noticed that 693 Mr was higher on steeper slopes (Fig. B3). Contrariwise, root biomass tends to be distributed 694 towards deeper portions of the soil when there is more SOC, which would reduce the amount 695 of root biomass allocated near the surface (Fig. B3) and thus to Aro. All these aspects support 696 the assumption of using a 'cone-shape-volume' to model the distribution of root biomass in the

697 soil (Fig. 3b; Supplementary Material), which was further supported by the strong influence of 698 Mr on Aro (Figs.6b and 7). However, the observed mismatch between empirical and parametric 699 ecohydrological VRD models (Fig. 7a-c; Table 4), in which Aro, Mr, and b are embedded 700 (Table 1), suggests that the modelling approach to estimate the key VRD parameters Aro and 701 b (Eqs. 4, 5 and 7; Table 1) must be revised for alpine ecosystems. Future work may consider 702 to explore the effect of topography in detail and/or to include climate-specific variables such 703 as snowpack depth, duration of the snow-free period, and frozen ground cycle, and how these 704 influence AWP and Aro in alpine ecosystems. In addition, soil nutrient limitations (e.g. 705 nitrogen; Zong et al., 2020) and plant-specific aspects related to growth and survival strategies 706 of alpine plants (e.g. Cooper et al., 2003; Germino, 2014) can be considered in future versions 707 of the VRD model to portray more realistically the characteristic allocation of plant biomass 708 above- and belowground in alpine ecosystems (e.g. Wu et al., 2013).

709

710 Nonetheless, the strong correlation between the investigated plant attributes and the fitting 711 parameters resulting from evaluating the relationship between empirical and parametric, 712 ecohydrological VRD models (Table 6) suggests that the collection of studied plant attributes 713 was appropriate, setting the direction of future research. The strong influence of plant attributes 714 such as the crown spread and projected area on the plant aboveground biomass (Ma) and Aro 715 (Figs.6b-c, 7, Appendix B – Figs. B2 and B3) hints at the possibility of establishing robust 716 data-mining approaches able to predict VRD on the basis of easy-to-measure aboveground 717 plant attributes (e.g. Fig. 3); provided that information on the allometry relationship between 718 above and belowground plant parts is available (Table 2; e.g. Cheng and Niklas, 2007). In this 719 regard, we expected that the allometry ratio (i.e. ALR=Mr/Ma; Table 2) would be consistently 720 above unity in the studied alpine plant species (e.g. Pohl et al., 2011), as a greater allocation of biomass to the belowground plant parts could help plants to withstand harsh aboveground 721

conditions (Germino, 2014). Nonetheless, only *P. alpina* had an *ALR* consistently above unity.
Future work helping to establish consistent and site-specific allometry relationships between
above and belowground plant parts in alpine ecosystems (e.g. Štastná et al. 2012) will
undoubtedly help to consolidate approaches seeking to describe VRD using very few, easily
measurable, parameters, like the one discussed herein.

727

728 5. Conclusion

729 Our study consolidates a simple protocol to describe the vertical root distribution (VRD) in 730 herbaceous plants. It also addresses, for the first time, the influence of soil and plant attributes 731 on key VRD parameters, validating the principles and assumptions behind the existing 732 parametric, ecohydrological models predicting VRD, and casting light on how VRD can be 733 effectively described using simple climate, soil and plant attributes. In fact, we confirmed that 734 insights into the water mass balance in the soil and into the water available to plants are crucial 735 to describe VRD in alpine ecosystems, as it has been suggested in previous studies for semi-736 arid and temperate humid ecosystems. However, the existing parametric ecohydrological VRD 737 models were not able to portray successfully the vertical root distribution of the studied alpine 738 plants in the light of the measured root profiles. Although we found a strong correlation 739 between empirical and parametric VRD models that establish a clear direction for future 740 research, we also think that the parametric VRD model needs to be revised in the future to 741 include features affecting the water available to plants in alpine ecosystems, such as the 742 snowpack characteristics or the length of the snow-free and frozen ground periods. We also 743 encourage future work exploring in detail the effect of topography, elevation, climate and 744 nutrient limitations on VRD, as these factors can help to formulate new models predicting VRD 745 realistically.

747	CRediT authorship contribution statement			
748	Alejandro Gonzalez-Ollauri: funding acquisition, conceptualization, methodology, data			
749	curation, formal analysis, investigation, project administration, writing - original draft, writing			
750	– review & editing			
751	Csilla Hudek: investigation, writing – review & editing			
752	Slododan B. Mickovski: funding acquisition, methodology, project administration, writing -			
753	review & editing			
754	Davide Viglietti: investigation			
755	Nicole Ceretto: investigation			
756	Michele Freppaz: project administration, resources, writing – review & editing			
757				
758	Acknowledgement			
759				
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767	6. References			
768				
769	Alvarez-Uria, P., Körner, C., 2007. Low temperature limits of root growth in decidious and			
770	evergreen temperate tree species. Funct. Ecol. 21, 211–218.			

- Arnone, E., Caracciolo, D., Noto, L. V., Preti, F., and Bras, R. L., 2016. Modeling the
- hydrological and mechanical effect of roots on shallow landslides, *Water Resour*.
- 773 *Res.*, 52, 8590–8612, doi:<u>10.1002/2015WR018227</u>.
- ASTM (1995) D 4972-95a: Standard test method for pH of soils. ASTM International, West
- 775 Conshohocken, PA, USA.
- Asturnauta, 2020. *Euphrasia minima*. <u>https://www.asturnatura.com/especie/euphrasia-</u>
  minima.html. Retrieved on 07/07/2020.
- 778 Bengough A.G. (2003) Root Growth and Function in Relation to Soil Structure,
- 779 Composition, and Strength. In: de Kroon H., Visser E.J.W. (eds) Root Ecology.
- 780 Ecological Studies (Analysis and Synthesis), vol 168. Springer, Berlin, Heidelberg.
- 781 https://doi.org/10.1007/978-3-662-09784-7\_6
- 782 Blozan, W., 2008. Tree measuring guidelines of the eastern native tree society. Bull. East.
- 783 Native Tree Soc. 1 (1), 3–10.
- 784 Böhm, W., 1979. Methods of Studying Root Systems. Springer Verlag, Berlin.
- 785 Breiman, L., 2001. Random forests. Mach. Learn. 45, 5–32.
- Burylo, M., Hudek, C., Rey, F., 2011. Soil reinforcement by the roots of six dominant species
- 787 on eroded mountainous marly slopes (Southern Alps, France). Catena, 84: 70-78.
- 788 Burylo, M., Dutoit, T. and Rey, F., 2014. Species Traits as Practical Tools for Ecological
- 789 Restoration of Marly Eroded Lands. Restor Ecol, 22: 633-640. doi:<u>10.1111/rec.12113</u>
- 790 Casper, B.B., Schenk, H.J., Jackson, R.B., 2003. Defining a Plant's Belowground Zone of
- 791 Influence. Ecology 84 (9), 2313–2321.

- Cheng, D., Niklas, K.J., 2007. Above- and below-ground biomass relationships across 1534
  forested communities. Ann. Bot. 99, 95–102.
- Chiatante, D., Scippa, S.G., Di Lorio, A., Sarnataro, M., 2003. The influence of steep slopes
  on root system development. J. Plant Growth Regul. 21, 247–260.
- 796 Clothier, B.E., Green, S.R. and Deurer, M., 2008. Preferential flow and transport in soil:
- progress and prognosis. European Journal of Soil Science, 59: 2-13. doi:<u>10.1111/j.1365-</u>
  2389.2007.00991.x
- 799 Cooper, E., Dullinger, S., Semenchuk, P., 2011. Late snowmelt delays plant development and
- 800 results in lower reproductive success in the High Arctic. Plant science, 180:157-67.
- 801 10.1016/j.plantsci.2010.09.005.
- 802 Coutts, M.P., Nielsen, C.C.N., Nicoll, B.C., 1999. The development of symmetry, rigidity and
  803 anchorage in the structural root system of conifers. Plant and Soil 217, 1-15.
- 804 Darwin, C., 1880. The Power of Movement in Plants. John Murray, London, UK.
- 805 Fitter, A.H., Stickland, T.R., 1991. Architectural analysis of plant root systems 2. Influence of
- 806 nutrient supply on architecture in contrasting plant species. New phytologist, 118: 383-
- 807 389.
- 808 Forde, B., Lorenzo, H., 2001. The nutritional control of root development. Plant and
- 809 Soil, 232: 51–68. https://doi.org/10.1023/A:1010329902165
- 810 Freppaz, M., Filippa, G., Caimi, A., et al., 2010. Soil and plant characteristics in the alpine
- 811 tundra (NW Italy). In: Tundras: Vegetation, Wildlife and Climate Trends. Nova Publishers.
- 812 pp 81–110.

- 813 Freppaz, M., Viglietti, D., Balestrini, R., Lonati, M., Colombo, N., 2019. Climatic and
- 814 pedoclimatic factors driving C and N dynamics in soil and surface water in the alpine tundra
- 815 (NW-Italian Alps). Nature Conservation, 34: 67-90.
- Fuller, W. A. 1987. Measurement Error Models. John Wiley & Sons. ISBN 978-0-471-861874.
- 818 Gale J. (2004). Plants and altitude--revisited. Annals of botany, 94(2), 199.
  819 https://doi.org/10.1093/aob/mch143
- 820 Germino M.J. 2014. Plants in Alpine Environments. In: Monson R. (eds) Ecology and the
- 821 Environment. Springer, New York, NY
- 822 Goodman, A.M., Ennos, A.R., 1999. The Effects of Soil Bulk Density on the Morphology and
- 823 Anchorage Mechanics of the Root Systems of Sunfower and Maize. Annals of Botany, 83:

824 293-302.

- 825 Gonzalez-Ollauri, A., Mickovski, S.B., 2016. Using the root spread information of pioneer
- plants to quantify their mitigation potential against shallow landslides and erosion. Ecol. Eng.
  95, 302–315.
- 828 Gonzalez-Ollauri, A., Mickovski, S.B., 2017a. Plant-soil reinforcement response under
- 829 different soil hydrological regimes. Geoderma 285, 141–150.
- 830 Gonzalez-Ollauri, A. and Mickovski, S.B., 2017b. Plant-Best: A novel plant selection tool for
- 831 slope protection. Ecological Engineering 106 (2017) 154–173
- 832 Gonzalez-Ollauri, A. and Mickovski, S.B., 2017c. Shallow landslides as drivers for slope
- 833 ecosystems evolution and biophysical diversity. Landslides, 14(5), 1699-1714. DOI
  834 10.1007/s10346-017-0822-y
- 835 Gonzalez-Ollauri, A., Mickovski, S.B., 2018. Green for Brown (G4B): A Novel Tool for
- 836 Evaluating Phytoextraction in Soils Polluted by Heavy Metals. In: Kallel A., Ksibi M., Ben

- Bia H., Khélifi N. (eds) Recent Advances in Environmental Science from the EuroMediterranean and Surrounding Regions. EMCEI 2017. Advances in Science, Technology &
  Innovation (IEREK Interdisciplinary Series for Sustainable Development). Springer, Cham.
- 840 http://doi-org-443.webvpn.fjmu.edu.cn/10.1007/978-3-319-70548-4\_81
- 841 Gonzalez-Ollauri, A., Stokes, A., Mickovski, S.B., 2020. A novel framework to study the effect
- 842 of tree architecture in stemflow yield and its consequences for soil-water dynamics. Journal of
- 843 Hydrology, 582: 124448.
- 844 Greenwell, B.M., 2017. pdp: An R package for constructing partial dependence plots. R J. 9
  845 (1), 421–436.
- 846 Greve, P., Roderick, M.L., Ukkola, A.M., Wada, Y., 2019. The aridity index under global
  847 warming. Environ. Res. Lett., 14: 124006.
- Hastie, T., Tibshirani, R., Friedman, J., 2009. The Elements of Statistical Learning, Second
  Edition, Section 10.13.2, Springer, 2009.
- 850 Head, K.H., 1980. Manual of Soil Laboratory Testing. CRC Press, Boca Raton, US.
- 851 Herbert, D.A., Rastetter, E.B., Gough, L., Shaver, G.R., 2004. Species diversity across
- nutrient gradients: an analysis of resource competition in model ecosystems. Ecosystems
  7:296–310
- Hudek, C., Stanchi, S., D'Amico, M., Freppaz, M., 2017a. Quanitfying the contribution of the
- root system of alpine vegetation in the soil aggregate stability of moraine. International Soiland Water Conservation Research, 5: 36-42.
- 857 Hudek, C., Sturrock, C.J., Atkinson, B.S., Stanchi, S., Freppaz, M., 2017b. Root morphology
- and biomechanical characteristics of high-altitude alpine plant species and their potential in
- soil stabilization. Ecological Engineering, 109, Part B: 228-239.

- 860 Iversen, C. M., Sloan, V.L., Sullivan, P.F., Euskirchen, E.S., McGuire, A.D., Norby, R.J.,
- 861 Walker, A.P., Warren, J.M., Wullschleger, S.D., 2014. The unseen iceberg: plant roots in arctic
- tundra. New Phytologist, 205: 34-58.
- Jackson, R.B., Canadell, J., Ehleringer, J.R., Mooney, H.A., Sala, O.E., Schulze, E.D., 1996.
- A global analysis of root distributions for terrestial biomes. Oecologia 108, 389–411.
- Jarvis, N., 1989. A simple empirical model of root water uptake. Journal of Hydrology,
  107:57-72.
- 867 Kaspar, T.C., Bland, W. L., 1992. Soil temperature and root growth. Soil Science,

868 154(4):290-299.

- 869 Kirkham, M. B., 2005. Principles of Soil and Water Retentions. Elsevier, Amsterdam, NL.
- Kleidon, A., 2004. Global datasets of rooting zone depth inferred from inverse methods, J.
  Clim., 17(13), 2714–2722.
- Kokutse, N. K., Tranquille Temgoua, A. G., Kavazović, Z., 2016. Slope stability and
  vegetation: Conceptual and numerical investigation of mechanical effects. Ecological
  Engineering, 86:146-153, https://doi.org/10.1016/j.ecoleng.2015.11.005.
- Köppen, W., 1884. The thermal zones of the Earth according to the duration of hot, moderate
- amd cold periods and the impact of heat on the organic world. Meteorol. Z. 1, 215–226.
- Köstler, J. N., Bru ckner, E., et Bibelriether, H. 1968: Die Wurzeln der Waldbau me edn.
- 878 Verlag Paul Parey.
- Kuhn, M. et al., 2018. caret: Classification and Regression Training. R package version 6.
- 880 0.81. https://CRAN.R-project.org/package=caret.

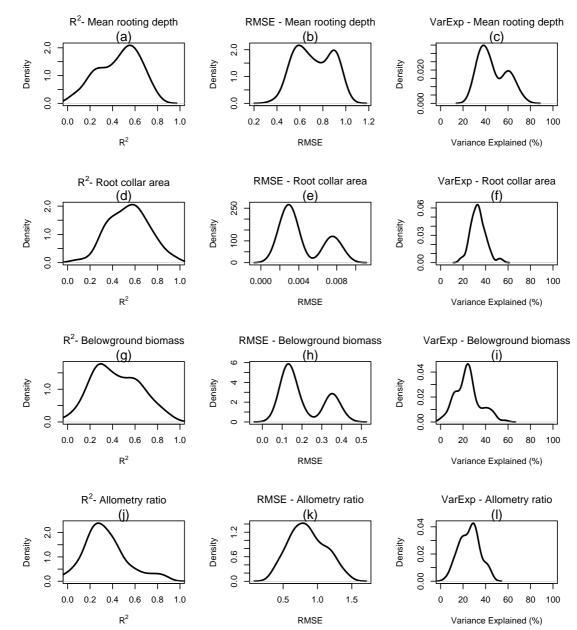
- 881 Kutschera L., Lichtenegger E., 1992. Wurzelatlas mitteleuropaischer Griinlandpflanzen. Band
- 2: Pteridophyta und Dicotyledoneae; Teil 1: Morphologie, Anatomie, Okologie, Verbreitung,
- 883 Soziologie, Wirtschaft. Gustav Fischer, Stuttgart
- Laio, F., 2006. A vertically extended stochastic model of soil moisture in the root zone. Water
  Resources Research, 42:W02406
- Laio, F., D'Odorico, P., Ridolfi, L., 2006. An analytical model to relate the vertical root
  distribution to climate and soil properties. Geophys. Res. Lett., 33 (2006), p. L18401
- Lahti, M., Aphalo, P., Finér, L., Ryyppö, A, Lehto, T., Mannerkoski, H., 2005. Effects of soil
- temperature on shoot and root growth and nutrient uptake of 5-year-old Norway spruce
  seedlings. Tree physiology, 25: 115-22. 10.1093/treephys/25.1.115.
- Liaw, A., Wiener, M., 2002. Classification and Regression by randomForest. R News 2(3):
- 892 18-22.
- Lu, N., Likos, W.J., 2004. Unsaturated Soil Mechanics. John Wiley and Sons, Hoboken,
- 894 US.
- 895 Lucherini, A., Gonzalez-Ollauri, A., Mickovski, S.B., 2020. The effect of vegetation on soil
- 896 polluted with galligu: phytostabilisation and novel approaches to evaluate soil galligu
- 897 concentration. Environmental Geotechnics, <u>https://doi.org/10.1680/jenge.19.00031</u>
- Lynch, J., 1995. Root architecture and plant productivity. Plant Physiol. 109:7-13.
- 899 Matthies, D. 1998. Influence of the host on growth and biomass allocation in the two facultative
- 900 root hemiparasites Odontites vulgaris and Euphrasia minima. Flora (1998) 193, 187-193
- 901 McMaster, G.S., Wilhelm, W.W., 1997. Growing degree-days: one equation, two
- 902 interpretations. Agric. Forest Meteorol. 87, 291–300.

- 903 Mickovski, S.B., van Beek, L.P.H. & Salin, F. Uprooting of Vetiver Uprooting Resistance of
- 904 Vetiver Grass (Vetiveria zizanioides). Plant Soil 278, 33–41 (2005).
- 905 Miyamoto, K., Wagai, R., Aiba, S. et al. Variation in the aboveground stand structure and
- 906 fine-root biomass of Bornean heath (kerangas) forests in relation to altitude and soil nitrogen
- 907 availability. Trees 30, 385–394 (2016). https://doi.org/10.1007/s00468-015-1210-7
- 908 Molotch, N.P., Brooks, P.D., Burns, S.P., Litvak, M., Monson, R.K., McConnell, J.R. and
- 909 Musselman, K. (2009), Ecohydrological controls on snowmelt partitioning in mixed-conifer
- 910 sub-alpine forests. Ecohydrol., 2: 129-142. doi:<u>10.1002/eco.48</u>
- 911 Pierce, S., Stirling, C. M., Baxter, R., 2000. Architectural and physiological heterogeneity
- 912 within the synflorescence of the pseudoviviparous grass *Poa alpina* var. vivipara L. Journal of
- 913 Experimental Botany, 51(351):1705-1712.
- Pohl, M., Stroude, R., Buttler, A., Rixen, C., 2011. Functional trails and root morphology of
- alpine plants. Annals of Botany, 108: 537-545.
- 916 Preti, F., Giadrossich, F., 2009. Root reinforcement and slope bioengineering sta-bilization
- 917 by Spanish Broom (Spartium junceum L.). Hydrol. Earth Syst. Sci. 13, 1713–1726.
- 918 Preti, F., Dani, A., Laio, F., 2010. Root profile assessment by means of hydrological,
- 919 pedological and above-ground vegetation information for bio-engineering purposes.
- 920 Ecological Engineering, 36:305-316.
- 921 Preti, F., 2013. Forest protection and protection forest: Tree root degradation over hydrological
- shallow landslides triggering. Ecological Engineering, 61P:633-645.
- Priestley, C., Taylor, R., 1972. On the assessment of surface heat flux and evaporation using
  large-Scale parameters. Mon. Weather Rev. 100 (2), 81–92.
- 925 Quine, C.P., Burnand, A.C., Coutts, M.P., Reynard, B.R., 1991. Effects of mounds and

- stumps on the root architecture of Sitka spruce on a peaty gley restocking site. Forestry 64,385-401.
- 928 R Core Team, 2018. R: A Language and Environment for Statistical Computing. R
- 929 Foundation for Statistical Computing, Vienna, Austria https://www.R-project.org/.
- 930 Rodríguez-Iturbe, I., & Porporato, A., 2005. Ecohydrology of Water-Controlled Ecosystems:
- 931 Soil Moisture and Plant Dynamics. Cambridge: Cambridge University Press.
- 932 doi:10.1017/CBO9780511535727
- 933 Schenk, H.J., Jackson, R.B., 2005. Mapping the global distribution of deep roots in relation to
- 934 climate and soil characteristics. Geoderma 126, 129–140.
- 935 Schenk, H., 2005. Vertical vegetation structure below ground: scaling from root to globe.
  936 Progr. Bot. 66, 341–373.
- 937 Shukla M. (2014) Soil Physics: An Introduction. Boca Raton, Florida: CRC Press.
- 938 Šťastná, P., Klimešová, J., Doležal, J., 2012. Altitudinal changes in the growth and allometry
- 939 of *Rumex alpinus*. Alp Botany 122, 35–44. https://doi.org/10.1007/s00035-012-0099-7
- 940
- 941 Stokes, A., Atger, C., Bengough, A.G., Fourcaud, T., Sidle, R.C., 2009. Desirable plant root
- 942 traits for protecting natural and engineered slopes against landslides. Plant and Soil 324 (1), 1–
- 943 30.
- 944 Strobl, C., Boulesteix, A., Kneib, T. et al., 2008. Conditional variable importance for random
- 945 forests. BMC Bioinformatics 9, 307. https://doi.org/10.1186/1471-2105-9-307
- 946 Tardio, G., Gonzalez-Ollauri, A., Mickovski, S.B., 2016. A non-invasive root distribution
- analysis methodology from a slope stability approach. Ecol. Eng. 97, 46–57.

- Taub, D.R., Goldberg, D., 1996. Root system topology of plants form habitats differing in soil
  resource availability. Functional Ecology 10, 258-264.
- Taylor, J. R., 1997. An introduction to error analysis: The study of uncertainties physical
  measurements (2<sup>nd</sup> Edition). University Science Books, CA, USA.
- 952 Toth, B., Weynants, M., Nemes, A., Mako, A., Bilas, G., Toth, G., 2015. New generation of
- 953 hydraulic pedotransfer functions for Europe. Eur. J. Soil Sci. 66, 226–238.
- 954 Tron, S., Dani, A., Laio, F., Preti, F., Ridolfi, L., 2014. Mean root depth estimation at
- landslide slopes. Ecol. Engine 69, 118–125.
- 956 Tsutsumi, D., Kosugi, K., Mizuyama, T., 2003. Effect of hydrotropism on root system
- development in soybean (Glycine max): growth experiments and model simulation. J. PlantGrowth Regul. 21, 441–458.
- 959Ukwildflowers(2020).Lucanthemopsisalpina.960https://www.ukwildflowers.com/Web\_pages/leucanthemopsis\_alpina\_alpine\_moon\_daisy.ht961m). Retrieved on 07/07/2020.
- van Wijk, M. T., and Bouten, W., 2001). Towards understanding tree root profiles: Simulating
- 963 hydrologically optimal strategies for root distribution, Hydrol. Earth Syst. Sci., 5(4), 629–644.
- Verma, P, George, KV, Singh, HV, Singh, SK, Juwarkar, A, Singh, RN, 2006. Modeling
  rhizofiltration: heavy-metal uptake by plant roots. Environmental Modeling and Assessment,
- 966 11, 387-394.
- Waisel, Y., Eshel, A., Kafkafi. U., (Eds.) 2002. Plant Roots: The Hidden Half, Marcel Dekker,
  New York.

- 969 Wu, J., Shen, Z., Zhang, X. et al. Biomass allocation patterns of alpine grassland species and
- 970 functional groups along a precipitation gradient on the Northern Tibetan Plateau. J. Mt. Sci.
- 971 10, 1097–1108 (2013). https://doi.org/10.1007/s11629-013-2435-9
- 2 Zeng, X. 2001. Global Vegetation Root Distribution for Land Modeling. Journal of
  3 Hydrometeorology, 2 (2001): 525-530.
- 274 Zhang J, Wang J, Chen J, Song H, Li S, Zhao Y, Tao J and Liu J (2019) Soil Moisture
- 975 Determines Horizontal and Vertical Root Extension in the Perennial Grass Lolium perenne L.
- 976 Growing in Karst Soil. Front. Plant Sci. 10:629. doi: 10.3389/fpls.2019.00629
- 2007 Zong, N, Song, M, Zhao, G, Shi, P., 2020. Nitrogen economy of alpine plants on the north
- 978 Tibetan Plateau: Nitrogen conservation by resorption rather than open sources through
- 979 biological symbiotic fixation. Ecol Evol. 2020; 10: 2051– 2061.
  980 https://doi.org/10.1002/ece3.6038
- Zuo, C., and R. Zhang (2002), Estimating root-water-uptake using and inverse method, Soil
  Sci., 167(9), 561–571.
- 983



# 984 Appendix A: Goodness of fit of the Random Forest models fitted to key vertical root

## 985 distribution parameters



Figure A1. Probability density functions illustrating the coefficients of determination (R<sup>2</sup>), root mean square error (RMSE) and
 variance explained (VarExp) retrieved from the cross-validation process implemented on random forest (RF) models, fitted
 between the studied soil and plant attributes to predict key vertical root distribution parameters.

Table A1. Coefficient of determination ( $R^2$ ) and root mean square error (RMSE) for the best performing random forest

994 995 996 models fitted between the key vertical root distribution (VRD) parameters mean rooting depth (b), cross-sectional area at the root collar (Aro) and root biomass (Mr) and the studied plant and soil attributes.

	R <sup>2</sup>	RMSE	Model No.
b	0.78	0.67	28
Aro	0.92	0.003	78
Mr	0.90	0.11	68

997

### 999 Appendix B: Partial dependence plots (PDPs) for the key vertical root distribution

1000 parameters

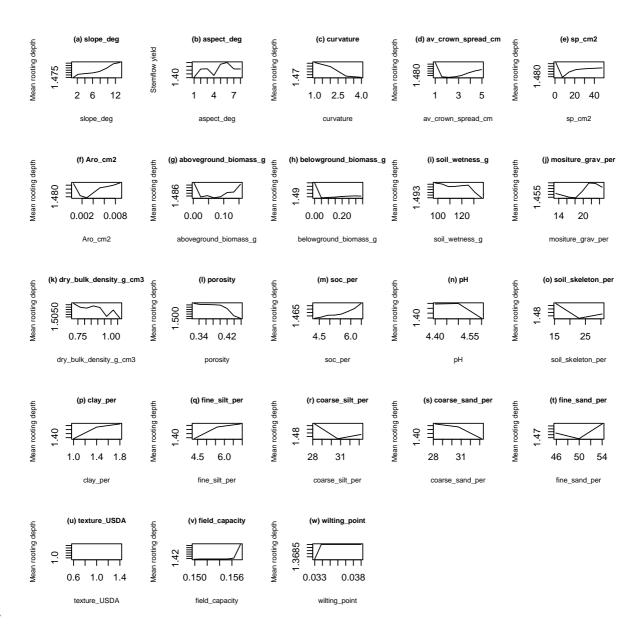
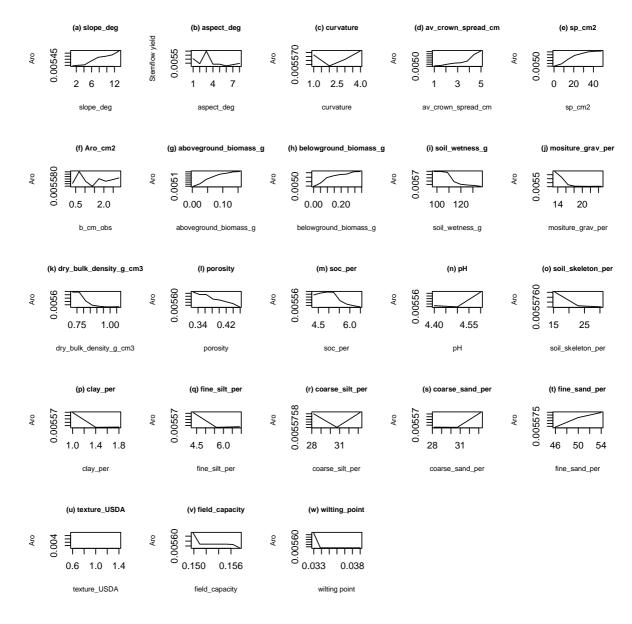
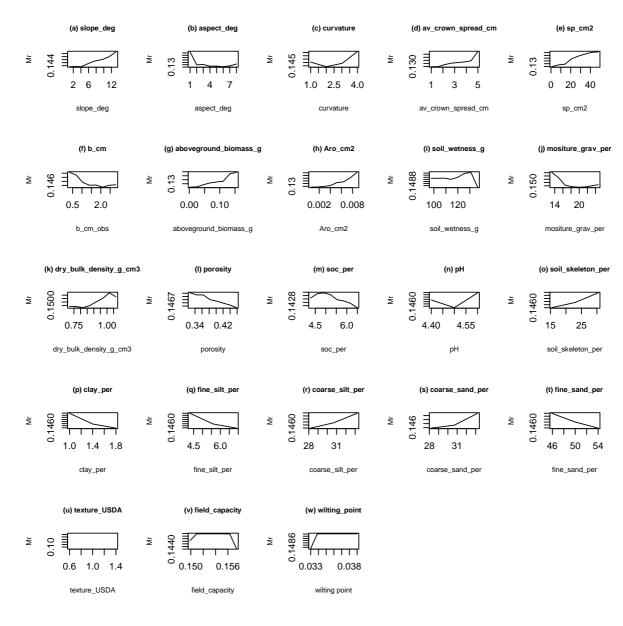


Figure B1. Partial dependence plots (PDPs) showing the relationship between the mean rooting depth (b) and the investigated
 plant and soil attributes in this study retrieved from fitting random forest models as indicated in the analytical framework
 shown in Figure 4.



1010 Figure B2. Partial dependence plots (PDPs) showing the relationship between the cross-sectional area at the root collar (Aro) and the investigated plant and soil attributes in this study retrieved from fitting random forest models as indicated in the

- 1012 analytical framework shown in Figure 4.



1020 Figure B3. Partial dependence plots (PDPs) showing the relationship between the root biomass (Mr) and the investigated

1021 plant and soil attributes in this study retrieved from fitting random forest models as indicated in the analytical framework shown Figure 4.

### 1024 Appendix C. R script – VRD data mining

```
1025 \\ 1026 \\ 1027 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 1028 \\ 
                    #Description:
                    #this script provides a series of functions to evaluate the relationship between the 'rooting
                      depth' variable and selected soil and plant attributes. Please, note that this is just a
                      sample script for the 'rooting depth' only.
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1038
1039
                    #Instructions:
                    #copy-paste the following script in your R console https://cran.r-project.org/ and change the
                      working directory as appropriate
                    ******
                    #Outline
                    #########
                    #i-RANDOM FOREST IMPLEMENTATION: it fits random forest models (RF) between the target
                      variable and selected plant and soil attributes and generates data frames storing relevant
                      outputs related to goodness of fit and relative importance of covariates
                    #ii-OUTPUTS COLLECTION AND RELATIVE IMPORTANCE EVALUATION: outputs collection, generation of
                     output datasets and creation of relevant plots evaluating model quality
1040
                    #iii-PARTIAL DEPENDENCE PLOTS: creates practical dependence plots for the target variable
1041
1042
1043
1044
                    ****************
                    #Abbreviations:
                    #b cm obs: measured mean rooting depth in cm
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1057
                    *****
                    #load data set and required R packages
                    setwd("/Users/ollauri/Desktop/work/catena roots/in") #write the path to your folder here
                    dts<-read.csv("DATASET.csv")
                    library(caret)
                    library(rattle)
                    library(pdp)
                    library(randomForest)
1058
1059
                    #i: RANDOM FOREST IMPLEMENTATION
                    1060
1061
1062
                    n < -100 #number of models to fit
1063
1064
                    RFs.b<-vector("list",n) #empty list object to store RF models</pre>
                    predictions.b<-vector("list", n) #empty list object to store predictions from RF models
 1065
                    train.A<-vector("list",n) #empty list object to store train data sets to fit RFs</pre>
 1066
                    LMs.b<-vector("list",n) #empty list object to store objective function -i.e. linear</pre>
1067 \\ 1068
                      regression models between observed and predicted
                    RMSEs.b<-vector("list",n) #empty list to store RMSE (root mean square error)
1069
                    Rsq.b<-vector("list",n) #empty list to store coefficient of determination (r-sq)
1070
                    out.b<-list() #empty list to store outputs</pre>
1071
1072
1073
1074
                    ct<-seq(1,100) #vector to assign numbers to outputs</pre>
                    output.b<-matrix(,nrow=n,ncol=3) #empty matrix to store outputs</pre>
                    varImp.b<-vector("list",n) #empty list to store variables importance</pre>
                    varImp.vals.b<-list() #empty list to store values of relative importance</pre>
1075
                    varImp.names.b<-list() #empty list to sotore variables names related to relative importance
 1075
1076
1077
1078
1079
                    for(i in 1:n) {
                      set.seed(i) #random number changes in each model run
                      train.A[[i]]<-sample(nrow(dts), 0.7*nrow(dts)) #sets training data set - i.e. bootstrapping</pre>
                      RFs.b[[i]]<-randomForest(b_cm_obs~species+site+slope_deg+aspect_deg+curvature+
av_crown_spread_cm+ sp_cm2+ Aro_cm2+aboveground_biomass_g+belowground_biomass_g+
1080
1081 \\ 1082
                      soil_wetness_g+ mositure_grav_per+ dry_bulk_density_g_cm3+ porosity+ soc_per+ pH+
soil_skeleton_per+ clay_per+ fine_silt_per+ coarse_silt_per+ coarse_sand_per+ fine_sand_per+
1083 \\ 1084 \\ 1085 
                      texture USDA+ field capacity+
                      wilting_point,data=dts[train.A[[i]],],mtry=5,importance=TRUE,ntree=1000) #fits random forest
                      models between target variable and soil & plant attributes
1086 \\ 1087 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 1088 \\ 
                      predictions.b[[i]]<-predict(RFs.b[[i]],dts[-train.A[[i]],]) #predictions using the RF models</pre>
                      and remaining data set
                      LMs.b[[i]]<-lm(predictions.b[[i]]~dts$b cm obs[-train.A[[i]]]) #fits regression models</pre>
 1089
1090
                      RMSEs.b[[i]]<-sqrt(mean((dts$b_cm_obs[-train.A[[i]]]-predictions.b[[i]])^2)) #calculates RMSE
Rsq.b[[i]]<-as.matrix(summary(LMs.b[[i]])$adj.r.squared) #calculates r-sq</pre>
  .091
                      out.b[[i]]<-list(ct[i],Rsq.b[[i]],RMSEs.b[[i]]) #stores outputs</pre>
   092
                      output.b[i,]<-c(out.b[[i]][[1]],as.numeric(out.b[[i]][[2]]),out.b[[i]][[3]]) #arranges</pre>
                      outputs
                      varImp.b[[i]]<-varImp(RFs.b[[i]]) #calculates RELATIVE IMPORTANCE</pre>
```

```
1095
                                      varImp.vals.b[[i]]<-varImp.b[[i]] #stores relative importance values</pre>
 1095
1096
1097
1098
1099
                                     varImp.names.b[[i]]<-rownames(varImp.b[[i]])#stores variables names</pre>
                                   }
1100
1101
1102
                                   #~~
                                   #ii: OUTPUTS COLLECTION
                                   #~~~
1103
1104
1104
1105
                                  setwd("/Users/ollauri/Desktop/work/catena roots/out")
                                  pdf("RFs_hist_rooting_depth_global.pdf")
hist(output.b[,2],col="gray",main="b vs plant &
    soil",xlab=expression(paste("R"^"2")),ylim=c(0,40)) #histogram showing density distribution
 1106
1107
                                      function for goodness of fit of RF models
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1115
                                  dev.off()
                                  out.df.b<-data.frame(model=output.b[,1],Rsq=output.b[,2],RMSE=output.b[,3])</pre>
                                  write.csv(out.df.b,"b global RFs.csv") #data frame storing summary for RF models' goodness of
                                    fit
                                  save(RFs.b,file="RFs b global.RData") #data base storing RF models
                                  varImp.df.b<-data.frame(var=unlist(varImp.names.b), imp=unlist(varImp.vals.b))</pre>
                                  write.csv(varImp.df.b, "varImp b.csv") #data frame storing relative influence
                                  pdf("RFs_b_varImp.pdf")
\begin{array}{c} 1116\\ 1117\\ 1118\\ 1119\\ 1120\\ 1121\\ 1122\\ 1123\\ 1124\\ 1125\\ 1126\\ 1127\\ 1128\\ 1132\\ 1133\\ 1133\\ 1133\\ 1135\\ 1137\\ 1138\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\ 1139\\
                                  boxplot(imp~reorder(var,imp,FUN=mean),data=varImp.df.b,horizontal=TRUE,las=2,cex.axis=0.7,mai
                                    n="varImp rooting depth") #boxplot for relative influence
                                  dev.off()
                                  #iii:PARTIAL DEPENDENCE PLOTS
                                   #Note: numerical variables, only - factor/character must be coded into numeric variables
                                      first
                                  #pl.x<-vector("list",n) #empty vectors to store partial dependence</pre>
                                  p2.x<-vector("list",n)
                                  p3.x<-vector("list",n)
                                   #p4.x<-vector("list",n)</pre>
                                  #p5.x<-vector("list",n)</pre>
                                 p6.x<-vector("list",n)
                                 p7.x<-vector("list",n)
    140
                                 p8.x<-vector("list",n)
 1141
1142
                                 p9.x<-vector("list"
                                 p10.x<-vector("list",n)</pre>
     142 \\ 143 \\ 144 \\ 145 \\ 146 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 \\ 147 
                                  pll.x<-vector("list",n)
                                  p12.x<-vector("list",n)
                                 p13.x<-vector("list",n)
                                  p14.x<-vector("list",n)
                                 p15.x<-vector("list",n)
 1148
1149
                                  pl6.x<-vector("list",n)</pre>
                                 p17.x<-vector("list",n)
   150
1151
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1159
                                 p18.x<-vector("list",n)
                                  p19.x<-vector("list",n)
                                  p20.x<-vector("list",n)
                                  p21.x<-vector("list",n)
                                  p22.x<-vector("list",n)
                                   #p23.x<-vector("list",n)</pre>
                                  p24.x<-vector("list",n)
                                  p25.x<-vector("list",n)
                                   #store partial dependence for each covariate with the target variable
     160
                                  for(i in 1:n) {
1161 \\ 1162 \\ 1163 \\ 1164 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 1165 \\ 
                                  #p1.x[[i]]<-partial(RFs.b[[i]],pred.var="species",plot=FALSE,from=0,to=10)</pre>
                                  p2.x[[i]]<-partial(RFs.b[[i]],pred.var="site",plot=FALSE)</pre>
                                  p3.x[[i]]<-partial(RFs.b[[i]],pred.var="slope deg",plot=FALSE)</pre>
                                   #p4.x[[i]]<-partial(RFs.b[[i]],pred.var="aspect_deg",plot=FALSE)</pre>
                                   #p5.x[[i]]<-partial(RFs.b[[i]],pred.var="curvature",plot=FALSE)</pre>
     165
     166
167
168
169
                                  p6.x[[i]]<-partial(RFs.b[[i]],pred.var="av_crown_spread_cm",plot=FALSE)</pre>
                                  p7.x[[i]]<-partial(RFs.b[[i]],pred.var="sp_cm2",plot=FALSE)
p8.x[[i]]<-partial(RFs.b[[i]],pred.var="Aro_cm2",plot=FALSE)</pre>
                                  p9.x[[i]]<-partial(RFs.b[[i]],pred.var="aboveground_biomass_g",plot=FALSE)
p10.x[[i]]<-partial(RFs.b[[i]],pred.var="belowground_biomass_g",plot=FALSE)</pre>
      170
 1171
                                  p11.x[[i]]<-partial(RFs.b[[i]],pred.var="soil wetness g",plot=FALSE)</pre>
```

```
\begin{array}{c} 1172\\ 1173\\ 1174\\ 1175\\ 1176\\ 1177\\ 1178\\ 1179\\ 1180\\ 1181\\ 1182\\ 1183\\ 1184\\ 1185\\ 1186\\ 1186\\ 1188\\ 1188\\ 1189\\ 1190\\ 1190\\ \end{array}
                                1191
1192
                                1193
1194
                                1195
                \begin{array}{c} 1196\\ 1197\\ 1198\\ 1197\\ 1198\\ 1199\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\ 1200\\
```

```
p12.x[[i]]<-partial(RFs.b[[i]],pred.var="mositure grav per",plot=FALSE)
p13.x[[i]]<-partial(RFs.b[[i]],pred.var="dry_bulk_density_g_cm3",plot=FALSE)
p14.x[[i]]<-partial(RFs.b[[i]],pred.var="porosity",plot=FALSE)</pre>
p15.x[[i]]<-partial(RFs.b[[i]],pred.var="soc per",plot=FALSE)
p16.x[[i]]<-partial(RFs.b[[i]],pred.var="pH",plot=FALSE)</pre>
p17.x[[i]]<-partial(RFs.b[[i]],pred.var="soil skeleton per",plot=FALSE)
p18.x[[i]]<-partial(RFs.b[[i]],pred.var="clay_per",plot=FALSE)
p19.x[[i]]<-partial(RFs.b[[i]],pred.var="fine silt per",plot=FALSE)
p20.x[[i]]<-partial(RFs.b[[i]],pred.var="coarse_silt_per",plot=FALSE)
p21.x[[i]]<-partial(RFs.b[[i]],pred.var="coarse_sand_per",plot=FALSE)
p22.x[[i]]<-partial(RFs.b[[i]],pred.var="fine_sand_per",plot=FALSE)</pre>
#p23.x[[i]]<-partial(RFs.b[[i]],pred.var="texture_USDA",plot=FALSE)</pre>
p24.x[[i]]<-partial(RFs.b[[i]],pred.var="field capacity",plot=FALSE)
p25.x[[i]]<-partial(RFs.b[[i]),pred.var="wilting point",plot=FALSE)</pre>
#retrieve outputs and arrange them for graphic display
#p1.a<-list()</pre>
#for(i in 1:n) {
#
       p1.a[[i]]<-p1.x[[i]][[1]]
#}
#p1.mtx.a<-do.call(rbind,p1.a)</pre>
#pl.mtx.a.t<-t(pl.mtx.a)</pre>
#p1.xs<-rowMeans(p1.mtx.a.t)</pre>
#pl.b<-list()</pre>
#for(i in 1:n) {
 #p1.b[[i]]<-p1.x[[i]][[2]]</pre>
#}
#pl.mtx<-do.call(rbind,pl.b) #i kind of have now the list elements in a matrix</pre>
#pl.mtx.t<-t(pl.mtx)</pre>
#p1.yhat<-rowMeans(p1.mtx.t)</pre>
p2.a<-list()
for(i in 1:n) {
p2.a[[i]]<-p2.x[[i]][[1]]
}
p2.mtx.a<-do.call(rbind,p2.a)
p2.mtx.a.t<-t(p2.mtx.a)
p2.xs<-rowMeans(p2.mtx.a.t)
p2.b<-list()
for(i in 1:n) {
p2.b[[i]]<-p2.x[[i]][[2]]
p2.mtx<-do.call(rbind,p2.b)
p2.mtx.t<-t(p2.mtx)
p2.yhat<-rowMeans(p2.mtx.t)
p3.a<-list()
for(i in 1:n) {
p3.a[[i]]<-p3.x[[i]][[1]]
-}
p3.mtx.a<-do.call(rbind,p3.a)
p3.mtx.a.t<-t(p3.mtx.a)
p3.xs<-rowMeans(p3.mtx.a.t)
p3.b<-list()
for(i in 1:n) {
p3.b[[i]]<-p3.x[[i]][[2]]
p3.mtx<-do.call(rbind,p3.b)
p3.mtx.t<-t(p3.mtx)
p3.yhat<-rowMeans(p3.mtx.t)
#p4.a<-list()</pre>
#for(i in 1:n) {
       p4.a[[i]]<-p4.x[[i]][[1]]
#
#}
#p4.mtx.a<-do.call(rbind,p4.a)</pre>
#p4.mtx.a.t<-t(p4.mtx.a)</pre>
#p4.xs<-rowMeans(p4.mtx.a.t)</pre>
#p4.b<-list()</pre>
#for(i in 1:n) {
       p4.b[[i]]<-p4.x[[i]][[2]]
#}
#p4.mtx<-do.call(rbind,p4.b)</pre>
#p4.mtx.t<-t(p4.mtx)</pre>
```

```
\begin{array}{l} 1249\\ 1250\\ 1252\\ 1253\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\ 1255\\
                         #p4.yhat<-rowMeans(p4.mtx.t)</pre>
                         #p5.a<-list()</pre>
                         #for(i in 1:n) {
                         #
                                          p5.a[[i]]<-p5.x[[i]][[1]]
                        #}
                        #p5.mtx.a<-do.call(rbind,p5.a)</pre>
                         #p5.mtx.a.t<-t(p5.mtx.a)</pre>
                         #p5.xs<-rowMeans(p5.mtx.a.t)</pre>
                         #p5.b<-list()</pre>
                         #for(i in 1:n) {
                                         p5.b[[i]]<-p5.x[[i]][[2]]
                         #
                         #}
                        #p5.mtx<-do.call(rbind,p5.b)</pre>
                        #p5.mtx.t<-t(p5.mtx)
                        #p5.yhat<-rowMeans(p5.mtx.t)</pre>
                        p6.a<-list()
                        for(i in 1:n) {
                          p6.a[[i]]<-p6.x[[i]][[1]]
                        p6.mtx.a<-do.call(rbind,p6.a)
                        p6.mtx.a.t<-t(p6.mtx.a)
                        p6.xs<-rowMeans(p6.mtx.a.t)
                        p6.b<-list()
                        for(i in 1:n) {
                         p6.b[[i]]<-p6.x[[i]][[2]]
                        }
                        p6.mtx<-do.call(rbind,p6.b)
                       p6.mtx.t<-t(p6.mtx)
                       p6.yhat<-rowMeans(p6.mtx.t)
                        p7.a<-list()
                        for(i in 1:n) {
                        p7.a[[i]]<-p7.x[[i]][[1]]
                        p7.mtx.a<-do.call(rbind,p7.a)
                        p7.mtx.a.t<-t(p7.mtx.a)
                        p7.xs<-rowMeans(p7.mtx.a.t)
                        p7.b<-list()
                        for(i in 1:n) {
                          p7.b[[i]]<-p7.x[[i]][[2]]
                        p7.mtx<-do.call(rbind,p7.b)
                        p7.mtx.t<-t(p7.mtx)
                        p7.yhat<-rowMeans(p7.mtx.t)
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13221
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13223
13225
                        p8.a<-list()</pre>
                        for(i in 1:n){
                          p8.a[[i]]<-p8.x[[i]][[1]]
                         }
                        p8.mtx.a<-do.call(rbind,p8.a)
                        p8.mtx.a.t<-t(p8.mtx.a)
                        p8.xs<-rowMeans(p8.mtx.a.t)
                        p8.b<-list()
                         for(i in 1:n) {
                          p8.b[[i]]<-p8.x[[i]][[2]]
                        p8.mtx<-do.call(rbind,p8.b)
                        p8.mtx.t<-t(p8.mtx)
                       p8.yhat<-rowMeans(p8.mtx.t)
                        p9.a<-list()</pre>
                        for(i in 1:n) {
                          p9.a[[i]]<-p9.x[[i]][[1]]
                        p9.mtx.a<-do.call(rbind,p9.a)
                        p9.mtx.a.t<-t(p9.mtx.a)
                        p9.xs<-rowMeans(p9.mtx.a.t)
                        p9.b<-list()
                        for(i in 1:n) {
                         p9.b[[i]]<-p9.x[[i]][[2]]
                        p9.mtx<-do.call(rbind,p9.b)
                        p9.mtx.t<-t(p9.mtx)
                        p9.yhat<-rowMeans(p9.mtx.t)
```

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          p10.a<-list()
          for(i in 1:n) {
           p10.a[[i]]<-p10.x[[i]][[1]]
          p10.mtx.a<-do.call(rbind,p10.a)
          pl0.mtx.a.t<-t(pl0.mtx.a)
          p10.xs<-rowMeans(p10.mtx.a.t)
          p10.b<-list()
          for(i in 1:n){
          p10.b[[i]]<-p10.x[[i]][[2]]
          p10.mtx<-do.call(rbind,p10.b)
          p10.mtx.t<-t(p10.mtx)
          p10.yhat<-rowMeans(p10.mtx.t)
          p11.a<-list()
          for(i in 1:n) {
          p11.a[[i]]<-p11.x[[i]][[1]]
          pl1.mtx.a<-do.call(rbind,pl1.a)</pre>
          pl1.mtx.a.t<-t(pl1.mtx.a)</pre>
          p11.xs<-rowMeans(p11.mtx.a.t)
          p11.b<-list()
          for(i in 1:n) {
p11.b[[i]]<-p11.x[[i]][[2]]
          pl1.mtx<-do.call(rbind,pl1.b)</pre>
          p11.mtx.t<-t(p11.mtx)
         pl1.yhat<-rowMeans(pl1.mtx.t)</pre>
          p12.a<-list()
          for(i in 1:n){
          p12.a[[i]]<-p12.x[[i]][[1]]
         p12.mtx.a<-do.call(rbind,p12.a)
          p12.mtx.a.t<-t(p12.mtx.a)
          p12.xs<-rowMeans(p12.mtx.a.t)
          p12.b<-list()
          for(i in 1:n) {
          p12.b[[i]]<-p12.x[[i]][[2]]
          p12.mtx<-do.call(rbind,p12.b)
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          p12.mtx.t<-t(p12.mtx)
          p12.yhat<-rowMeans(p12.mtx.t)
          p13.a<-list()
          for(i in 1:n) {
          p13.a[[i]]<-p13.x[[i]][[1]]
          p13.mtx.a<-do.call(rbind,p13.a)
          p13.mtx.a.t<-t(p13.mtx.a)
          p13.xs<-rowMeans(p13.mtx.a.t)
          p13.b<-list()
          for(i in 1:n) {
          p13.b[[i]]<-p13.x[[i]][[2]]
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         p13.mtx<-do.call(rbind,p13.b)
          p13.mtx.t<-t(p13.mtx)
         p13.yhat<-rowMeans(p13.mtx.t)
          p14.a<-list()
          for(i in 1:n) {
          p14.a[[i]]<-p14.x[[i]][[1]]
          p14.mtx.a<-do.call(rbind,p14.a)
          p14.mtx.a.t<-t(p14.mtx.a)
          p14.xs<-rowMeans(p14.mtx.a.t)
          p14.b<-list()
          for(i in 1:n) {
          p14.b[[i]]<-p14.x[[i]][[2]]
          p14.mtx<-do.call(rbind,p14.b)
          p14.mtx.t<-t(p14.mtx)
          p14.yhat<-rowMeans(p14.mtx.t)
 401
          p15.a<-list()
1402
          for(i in 1:n) {
```

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1403
                                       p15.a[[i]]<-p15.x[[i]][[1]]
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                                    p15.mtx.a<-do.call(rbind,p15.a)</pre>
                                    p15.mtx.a.t<-t(p15.mtx.a)
                                    p15.xs<-rowMeans(p15.mtx.a.t)
                                    _
p15.b<-list()
  1409
                                     for(i in 1:n) {
1409 \\ 1410 \\ 1411 \\ 1412 \\ 1413 \\ 1414 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 1415 \\ 
                                        p15.b[[i]]<-p15.x[[i]][[2]]
                                    p15.mtx<-do.call(rbind,p15.b)
                                    p15.mtx.t<-t(p15.mtx)
                                    p15.yhat <- rowMeans (p15.mtx.t)
\begin{array}{c} 1416\\ 1417\\ 1418\\ 1419\\ 1420\\ 1421\\ 1422\\ 1423\\ 1424\\ 1425\\ 1426\\ 1427\\ 1428\\ 1426\\ 1427\\ 1428\\ 1430\\ 1431\\ 1432\\ 1433\\ 1434\\ 1435\\ 1436\\ 1437\\ 1438\\ 1439\\ 1439\end{array}
                                    p16.a<-list()
                                     for(i in 1:n) {
                                      p16.a[[i]]<-p16.x[[i]][[1]]
                                    pl6.mtx.a<-do.call(rbind,pl6.a)</pre>
                                    p16.mtx.a.t<-t(p16.mtx.a)
                                    p16.xs<-rowMeans(p16.mtx.a.t)
                                    p16.b<-list()
                                     for(i in 1:n) {
                                       p16.b[[i]]<-p16.x[[i]][[2]]
                                    p16.mtx<-do.call(rbind,p16.b)
                                    p16.mtx.t<-t(p16.mtx)
                                   p16.yhat<-rowMeans(p16.mtx.t)
                                    p17.a<-list()
                                     for(i in 1:n) {
                                      p17.a[[i]]<-p17.x[[i]][[1]]
                                      3
                                    p17.mtx.a<-do.call(rbind,p17.a)
                                    p17.mtx.a.t<-t(p17.mtx.a)
                                    p17.xs<-rowMeans(p17.mtx.a.t)
                                    p17.b<-list()
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1444
                                      for(i in 1:n) {
                                       p17.b[[i]]<-p17.x[[i]][[2]]
                                    p17.mtx<-do.call(rbind,p17.b)
                                    p17.mtx.t<-t(p17.mtx)
1444 \\ 1445 \\ 1446 \\ 1447 \\ 1448 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1449 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 1440 \\ 
                                    p17.yhat<-rowMeans(p17.mtx.t)
                                    p18.a<-list()
                                     for(i in 1:n) {
                                       p18.a[[i]]<-p18.x[[i]][[1]]
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                                    p18.mtx.a<-do.call(rbind,p18.a)
                                    p18.mtx.a.t<-t(p18.mtx.a)
                                    p18.xs<-rowMeans(p18.mtx.a.t)
                                    p18.b<-list()
                                     for(i in 1:n) {
                                       p18.b[[i]]<-p18.x[[i]][[2]]
                                    p18.mtx<-do.call(rbind,p18.b)</pre>
                                    p18.mtx.t<-t(p18.mtx)
 1460
                                    p18.yhat<-rowMeans(p18.mtx.t)
1461
1462
1463
                                    p19.a<-list()
                                     for(i in 1:n) {
 1464
1465
                                      p19.a[[i]]<-p19.x[[i]][[1]]
                                      }
 146<u>6</u>
                                    p19.mtx.a<-do.call(rbind,p19.a)</pre>
 1467
                                    p19.mtx.a.t<-t(p19.mtx.a)
 1468
                                    p19.xs<-rowMeans(p19.mtx.a.t)
 1469
                                     p19.b<-list()
1470
1471
1472
1472
                                     for(i in 1:n) {
                                       p19.b[[i]]<-p19.x[[i]][[2]]
 1472
1473
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1476
1477
                                    p19.mtx<-do.call(rbind,p19.b)</pre>
                                    p19.mtx.t<-t(p19.mtx)
                                     p19.yhat<-rowMeans(p19.mtx.t)
                                     p20.a<-list()</pre>
     478
                                     for(i in 1:n) {
    479
                                        p20.a[[i]]<-p20.x[[i]][[1]]
```

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          p20.mtx.a<-do.call(rbind,p20.a)
          p20.mtx.a.t<-t(p20.mtx.a)
          p20.xs<-rowMeans(p20.mtx.a.t)
          p20.b<-list()
          for(i in 1:n) {
           p20.b[[i]]<-p20.x[[i]][[2]]
          p20.mtx<-do.call(rbind,p20.b)
          p20.mtx.t<-t(p20.mtx)
1490
          p20.yhat<-rowMeans(p20.mtx.t)
1491
1492
          p21.a<-list()
1493
          for(i in 1:n){
1494
1495
          p21.a[[i]]<-p21.x[[i]][[1]]
          }
1496
1496
1497
1498
          p21.mtx.a<-do.call(rbind,p21.a)
          p21.mtx.a.t<-t(p21.mtx.a)
          p21.xs<-rowMeans(p21.mtx.a.t)
1499
          p21.b<-list()</pre>
1500
1501
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1503
          for(i in 1:n) {
          p21.b[[i]]<-p21.x[[i]][[2]]
          p21.mtx<-do.call(rbind,p21.b)
15034

1505671508

1505671508

150671508

150671508

15071508

1511512

1511512

15115122234

155223155231153367

155321552234551

155332334567

15533678890

155423315556

15556

15556
          p21.mtx.t<-t(p21.mtx)
          p21.yhat<-rowMeans(p21.mtx.t)
          p22.a<-list()</pre>
          for(i in 1:n) {
          p22.a[[i]]<-p22.x[[i]][[1]]
          p22.mtx.a<-do.call(rbind,p22.a)
          p22.mtx.a.t<-t(p22.mtx.a)
          p22.xs<-rowMeans(p22.mtx.a.t)
          p22.b<-list()</pre>
          for(i in 1:n) {
           p22.b[[i]]<-p22.x[[i]][[2]]
          }
          p22.mtx<-do.call(rbind,p22.b)
          p22.mtx.t<-t(p22.mtx)
          p22.yhat<-rowMeans(p22.mtx.t)
          #p23.a<-list()</pre>
          #for(i in 1:n) {
                  p23.a[[i]]<-p23.x[[i]][[1]]
          #
          #}
          #p23.mtx.a<-do.call(rbind,p23.a)</pre>
          #p23.mtx.a.t<-t(p23.mtx.a)</pre>
          #p23.xs<-rowMeans(p23.mtx.a.t)</pre>
          #p23.b<-list()
          #for(i in 1:n) {
          #
                 p23.b[[i]]<-p23.x[[i]][[2]]
          #}
          #p23.mtx<-do.call(rbind,p23.b)</pre>
          #p23.mtx.t<-t(p23.mtx)</pre>
          #p23.yhat<-rowMeans(p23.mtx.t)</pre>
          p24.a<-list()
          for(i in 1:n){
           p24.a[[i]]<-p24.x[[i]][[1]]
          }
          p24.mtx.a<-do.call(rbind,p24.a)
          p24.mtx.a.t<-t(p24.mtx.a)
          p24.xs<-rowMeans(p24.mtx.a.t)
          p24.b<-list()
          for(i in 1:n) {
           p24.b[[i]]<-p24.x[[i]][[2]]
          1
          p24.mtx<-do.call(rbind,p24.b)
          p24.mtx.t<-t(p24.mtx)
          p24.yhat<-rowMeans(p24.mtx.t)
          p25.a<-list()
          for(i in 1:n) {
           p25.a[[i]]<-p25.x[[i]][[1]]
          p25.mtx.a<-do.call(rbind,p25.a)</pre>
```

}

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\begin{array}{c} 1557\\ 1558\\ 1559\\ 1560\\ 1561\\ 1562\\ 1563\\ 1563\\ 1564\\ 1565\end{array}
                        p25.mtx.a.t<-t(p25.mtx.a)
                        p25.xs<-rowMeans(p25.mtx.a.t)
                        p25.b<-list()
                         for(i in 1:n) {
                          p25.b[[i]]<-p25.x[[i]][[2]]
                        p25.mtx<-do.call(rbind,p25.b)</pre>
                        p25.mtx.t<-t(p25.mtx)
                        p25.yhat<-rowMeans(p25.mtx.t)
1566 \\ 1567 \\ 1568 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1569 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 1560 \\ 
                         #plot PDPs
1570
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                        pdf("RFs rooting depth PDPs.pdf")
                        par(mfrow=c(5,5))
                         #plot(p1.yhat[1:8]~p1.xs[1:8],type="l",xlab="species",ylab="Mean rooting depth",main="(a)
                          species",cex.main=0.8,cex.lab=0.8)
                        plot(p2.yhat[1:8]~p2.xs[1:8],type="l",xlab="site",ylab="Mean rooting depth",main="(b)
                          site",cex.main=0.8,cex.lab=0.8)
                        plot(p3.yhat[1:8]~p3.xs[1:8],type="l",xlab="slope deg",ylab="Mean rooting depth",main="(c)
                          slope deg",cex.main=0.8,cex.lab=0.8)
                         #plot(p4.yhat[1:8]~p4.xs[1:8],type="l",xlab="aspect deg",ylab="Mean rooting depth",main="(d)
                          aspect deg",cex.main=0.8,cex.lab=0.8)
                         #plot(p5.yhat[1:8]~p5.xs[1:8],type="1",xlab="curvature",ylab="Mean rooting depth",main="(e)
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                          curvature",cex.main=0.8,cex.lab=0.8)
                        plot(p6.yhat[1:8]~p6.xs[1:8],type="1",xlab="av crown spread cm",ylab="Mean rooting
                          depth",main="(f) av_crown_spread_cm",cex.main=0.8,cex.lab=0.8)
                        plot(p7.yhat[1:8]~p7.xs[1:8],type="l",xlab="sp_cm2",ylab="Mean rooting depth",main="(g)
sp_cm2",cex.main=0.8,cex.lab=0.8)
                        plot(p8.yhat[1:8]~p8.xs[1:8],type="l",xlab="Aro_cm2",ylab="Mean rooting depth",main="(h)
                          Aro cm2",cex.main=0.8,cex.lab=0.8)
                        plot (p9.yhat[1:8]~p9.xs[1:8],type="l",xlab="aboveground biomass g",ylab="Mean rooting
                        depth",main="(i) aboveground_biomass_g",cex.main=0.8,cex.lab=0.8)
plot(p10.yhat[1:8]~p10.xs[1:8],type="1",xlab="belowground_biomass_g",ylab="Mean rooting
depth",main="(j) belowground_biomass_g",cex.main=0.8,cex.lab=0.8)
plot(p11.yhat[1:8]~p11.xs[1:8],type="1",xlab="soil_wetness_g",ylab="Mean rooting
depth",main="(j) belowground_biomass_g",cex.main=0.8,cex.lab=0.8)
plot(p11.yhat[1:8]~p11.xs[1:8],type="1",xlab="soil_wetness_g",ylab="Mean rooting")

                          depth",main="(k) soil_wetness_g",cex.main=0.8,cex.lab=0.8)
                        plot(p12.yhat[1:8]~p12.xs[1:8],type="l",xlab="mositure grav per",ylab="Mean rooting
                          depth",main="(1) mositure_grav_per",cex.main=0.8,cex.lab=0.8)
                        depth / model__pit _pit _pit _pit _pit / model__pit _pit _pit (pi3.yhat[1:8]~p13.yhat[1:8], type="l", xlab="dry_bulk_density_g_cm3", ylab="Mean rooting
depth", main="(m) dry_bulk_density_g_cm3", cex.main=0.8, cex.lab=0.8)
plot(p14.yhat[1:8]~p14.xs[1:8], type="l", xlab="porosity", ylab="Mean rooting depth", main="(n)
 1599
                          porosity", cex.main=0.8, cex.lab=0.8)
1600
                        plot(p15.yhat[1:8]~p15.xs[1:8],type="l",xlab="soc per",ylab="Mean rooting depth",main="(o)
1601
1602
                          soc per", cex.main=0.8, cex.lab=0.8)
                        plot(p16.yhat[1:8]~p16.xs[1:8],type="l",xlab="pH",ylab="Mean rooting depth",main="(p)
1603
                          pH", cex.main=0.8, cex.lab=0.8)
1604
                        plot(p17.yhat[1:8]~p17.xs[1:8],type="1",xlab="soil skeleton per",ylab="Mean rooting
                        depth",main="(p) soil_skeleton_per",cex.main=0.8,cex.lab=0.8)
plot(p18.yhat[1:8]~p18.xs[1:8],type="l",xlab="clay_per",ylab="Mean rooting depth",main="(p)
clay_per",cex.main=0.8,cex.lab=0.8)
 1605
1606 \\ 1606 \\ 1607 \\ 1608 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1609 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 1000 \\ 
                        plot(p19.yhat[1:8]~p19.xs[1:8],type="1",xlab="fine silt per",ylab="Mean rooting
                          depth",main="(p) fine_silt_per",cex.main=0.8,cex.lab=0.8)
1610
1610
1611
1612
1613
                        plot(p20.yhat[1:8]~p21.xs[1:8],type="1",xlab="coarse_silt_per",ylab="Mean rooting
                        depth",main="(p) coarse_silt_per",cex.main=0.8,cex.lab=0.8)
plot(p21.yhat[1:8]~p21.xs[1:8],type="l",xlab="coarse_sand_per",ylab="Mean rooting
                          depth",main="(p) coarse_sand_per",cex.main=0.8,cex.lab=0.8)
                        plot(p22.yhat[1:8]~p22.xs[1:8],type="l",xlab="fine_sand_per",ylab="Mean rooting
depth",main="(p) fine_sand_per",cex.main=0.8,cex.lab=0.8)
1614
1615
                          #plot(p23.yhat[1:8]~p23.xs[1:8],type="1",xlab="texture USDA",ylab="Mean rooting
 1616
1617
1618
1619
1620
1621
1622
                          depth", main="(p) texture USDA", cex.main=0.8, cex.lab=0.8)
                        plot(p24.yhat[1:8]~p24.xs[1:8],type="1",xlab="field_capacity",ylab="Mean rooting
                          depth",main="(p) field_capacity",cex.main=0.8,cex.lab=0.8)
                         plot(p25.yhat[1:8]~p25.xs[1:8],type="l",xlab="wilting_point",ylab="Mean rooting
                          depth",main="(p) wilting point",cex.main=0.8,cex.lab=0.8)
                        dev.off()
1623
1624
```