

The influence of altitude on the distribution of subterranean organs and humus components in carpets of *Vaccinium myrtillus* (L.)

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Abstract. Humus profiles were sampled along an altitudinal gradient in the Macot-La-Plagne Forest (France, Northern Alps) in order to describe the variation occurring under carpets of bilberry [*Vaccinium myrtillus* (L.)] present within spruce forests. The vertical distribution of subterranean organs of bilberry (rhizomes and roots) was compared with i) that of spruce roots and other accompanying vegetation, ii) other components of humus profiles, in particular humified organic matter, mainly consisting of recent and old animal faeces. It was shown that bilberry roots were mostly concentrated in mineral horizons, while spruce roots and bilberry rhizomes rather occupied litter horizons. This was interpreted in terms of a strategy for capturing nutrients in the frame of the competition between spruce and bilberry. The effects of altitude were i) a change in the vegetation accompanying bilberry in dense bilberry carpets, bryophytes at the montane level being replaced by forbes at the subalpine level, ii) a decrease in the thickness of ectorganic horizons. This was interpreted as a shift from a moder system characterized by recalcitrant litter (moss) processed by an active faunal community (stabilized in the form of animal faeces), to a mor system characterized by low animal abundance but with litter of better quality which is easily leached in the absence of prominent faunal activity.

Keywords: Animal faeces, Bilberry, Humus form, Mountain, Rhizome, Root, Spruce

Nomenclature: Rameau et al. (1993) for plant species, Brêthes et al. (1995) for soil horizons.

Introduction

The age structure and the growth dynamics of bilberry (*Vaccinium myrtillus* L.) are well-known since the thorough examination of bilberry patches of varying age done by Flower-Ellis (1971) in Scotland. The rhizome of this acidophilic ericaceous shrub grows sympodially, forming clonal carpets which enlarge and densify in the course of time and accumulate raw humus (Bernier et al. 1993; Bernier & Ponge 1994; Maubon et al. 1995). The foliage of bilberry is deciduous and is richer in nutrients than that of most other Ericaceae, its soft leaf litter decaying easily despite a high tannin content (Gallet & Lebreton 1989; Gallet & Lebreton 1995). Given the nutritional demand of *V. myrtillus* (Ingestad 1973) we may wonder how it does not suffer from the shortage of nutrients resulting from podzolisation and the immobilization of nitrogen in the form of tannin-protein complexes (Wardle et al. 1997; Northup et al. 1998). The association of roots with mycorrhizal fungi able to use recalcitrant organic matter as a nitrogen source has been thought to explain the growth of ericaceous species in raw humus with a very low biological activity (Read & Kerley 1995; Näsholm et al. 1998). Nevertheless examination of the subterranean parts of bilberry reveals that raw humus is occupied by rhizomes while most roots grow more or less vertically through the mineral soil (Heath et al. 1938). Thus bilberry seems to exhibit a double strategy for capturing nutrients, by exploring the organic horizon with its rhizome system and the mineral soil with its root system.

In European mountains bilberry is commonly found in association with Norway spruce [*Picea abies* (L) Karst.], at least at the subalpine level (Gensac 1970). Recent studies revealed that Norway spruce and bilberry were in fact competing for the same micro-sites during their establishment, both germinating and growing better on mineral soils (André 1994; Ponge et al. 1998) and exhibiting biochemical-mediated antagonisms once established (Gallet 1994; Maubon et al. 1995; Jäderlund et al. 1996; Jäderlund et al. 1997). It was concluded that the development of bilberry as dense permanent carpets was favoured by thinning operations done in spruce forests, resulting in lack of spruce regeneration and thus in the long-term in a collapse of the forest ecosystem (Ponge et al. 1998). Within dense bilberry heath, spruce cannot establish successfully by seed. Conversely, bilberry declines when spruce crowns enlarge (Ponge et al. 1994; Maubon et al. 1995), possibly due to a combination of shading and nutrient depletion (Christy 1986). We may wonder how the antagonism

between bilberry and spruce is expressed when the subterranean parts of both species are present, as this is the case in dense bilberry carpets within bilberry-spruce forests, for instance by exploiting distinct horizons.

The thickness of organic matter accumulating on the forest floor in bilberry carpets has been observed to decrease with altitude (Ponge et al. 1998). This finding apparently contradicts the rule that more organic matter accumulates in the soil when the climate becomes colder and the substrate becomes poorer as occurs at higher elevation (Lichtenegger 1996; Körner 1999). The examination of the composition of organic matter (plant debris, animal faeces) might throw light on this unexpected phenomenon.

In order to answer the three above mentioned questions, it was decided to analyse the composition of humus profiles along an altitudinal gradient, in the Macot-La Plagne forest, where *V. myrtillus* and *P. abies* co-exist, dominating the forest/heath patchwork from the montane to the subalpine level (Gensac 1970; Ponge et al. 1994; Maubon et al. 1995). Micromorphological methods according to Ponge (1984), later on modified by Bernier & Ponge (1994), proved useful to quantify the composition of soil horizons at varying depths and compare humus profiles (Ponge 1999; Peltier et al. 2001). They were used in the present study, rather than washing techniques (McQueen 1968; Messier & Kimmins 1991; Dighton & Coleman 1992; Ehrenfeld et al. 1997), in order to estimate the abundance of fine roots. Together with dead and living plant parts, other components of humus profiles (animal faeces, mineral matter) will be quantified and compared from horizon to horizon and between humus profiles in order to characterize soil organic matter.

Study site

The Macot forest (Macot-la-Plagne, Savoy, France) is located on a north-exposed slope along the Tarentaise Valley, in the French northern Alps. The elevation ranges from 800m (near the Macot village) to 2100m (at the base of Mount Saint-Jacques). The substrate is poor in nutrients, arising from graywackes, schists and quartzites, with soils being acidic throughout. Due to a combination of favourable factors such as colluvial deposits and higher biological activity, the bottom of the slope is

characterized by richer soils. Thus, there is a gradient of increasing soil acidity with altitude (Loranger et al. 2001). Spruce is the dominant tree species, mixed with silver fir (*Abies alba* Mill.) at the montane level and with cembra pine (*Pinus cembra* L.) at the subalpine level, European larch (*Larix decidua* L.) being sparsely distributed over the whole altitudinal range. Bilberry is present at small isolated spots on rocky outcrops at the lower montane level, the size of carpets increasing with altitude, extending as pure ericaceous heath above the timberline, in admixture with other ericaceous species such as *Vaccinium vitis-idaea* L., *Rhododendron ferrugineum* L. and *Arctostaphylos uva-ursi* L. The higher montane and the lower subalpine levels are characterized by a mosaic assemblage of spruce forest and bilberry heath (the so-called bilberry-spruce forest) as a result of succession processes and silvicultural practices (Bernier & Ponge 1994; Ponge et al. 1994; Maubon et al. 1995; Ponge et al. 1998). Bilberry was often found associated with two mosses [*Rhytidiadelphus triquetus* (Hedw.) Warnst., *Hylocomium splendens* (Hedw.) B., S. & G.) and two grass species, the wavy hair-grass [*Avenella flexuosa* (L.) Parl.] and the greater wood-rush [*Luzula sylvatica* (Huds.) Gaud.].

Material and methods

Carpets of bilberry were sampled along a transect crossing the whole altitudinal range where bilberry was found in dense carpets. At 950, 1470, 1650, 1870 and 2150m a.s.l., after a cursory examination of physiognomic mosaics, one to three plots were selected representing the variety of bilberry carpets prevailing on the site. A total of 12 plots were thus selected for sampling humus profiles. Sites at 950 (2 samples) and 1470m (one sample) were the same as in Bernier & Ponge (1994) and Bernier (1996). Sites at 1650 (3 samples) and 1870m (3 samples) were the same as in Bernier et al. (1993). The site at 2150m (3 samples) was the same as in Ponge et al. (1994). At the centre of each plot a small humus block 5x5x15cm (lxwxh) was prepared with a sharp knife and the different layers were separated by hand and put immediately to small plastic jars filled with 95% ethyl alcohol.

The fixed material was examined under a dissecting microscope by pouring it, with as less disturbance as possible, in a Petri dish filled with ethyl alcohol. A transparent sheet with 600 points marked on it was put over the material and covered with alcohol, allowing for an estimation of the

volume of the different humus components in each layer. In order to increase the number of points, and thus the precision of the measurement, the grid was randomly displaced at the end of a counting run in order to allow for a new set of 600 points to be counted. This procedure was necessary for estimating the volume of very fine roots of bilberry. When no objects were visible under a point at the 40x magnification, the point was discarded. The total number of points taken into account for a given layer varied from 336 to 1603.

Plant organs were determined by help of a collection of main plant species growing in the vicinity of the sampled humus profiles: aerial and subterranean parts were fixed separately into ethyl alcohol and observed in the same conditions as humus components. Animal faeces and bodies were identified by morphological features (size, shape, colour, size of ingested particles) according to experience of the junior author.

Data were analysed by simple correspondence analysis (Greenacre 1984), which has been successfully applied to micromorphological data, allowing humus profiles as well as horizons to be compared and classified on the basis of their composition (Ponge 1999; Peltier et al. 2001). Humus components were used as active variables and layers of all humus profiles were used as observations. The five altitudes (950, 1470, 1870 and 2150m), the three arbitrary depth levels (O-5, 5-10 and 10-15cm) and the different horizons found (OL, OF, OH, A, E, B, S, rodent mound) were each used as passive variables. All variables were standardized, their mean being fixed to 20 and their variance to 1, for interpreting factorial coordinates as contributions to factorial axes (Ponge and Delhay 1995).

Given the absence of replication, no testing of hypothesis can be achieved on this data set, correspondence analysis being used only to reveal patterns hidden in a complex data matrix. Nevertheless, the significance of the first axis of correspondence analysis was tested by correlation analysis (Sokal & Rohlf 1995), as well as the effect of altitude on organic matter accumulation.

The mean depth of a humus component in a given profile was calculated using the vertical distribution of its percentage of occurrence. Thus each humus component can be characterized by an

array of twelve average depths, one for each profile. In turn, these average depths can be averaged, in order to give a global average depth (labeled mean depth) indicating the mean vertical position of a given humus component.

Results

One hundred and fifty-nine humus components were identified in the whole set of 53 samples (Table 1). A projection of humus components (active variables) and passive variables (elevation, depth level, horizon) in the plane of the first two axes of correspondence analysis (13% and 9% of total variance, respectively) revealed the influence of depth and altitude on the distribution of humus components and plant organs (Fig. 1). Axis 1 was correlated with depth (Fig. 2), thus it can be used to scale the different humus components according to their vertical distribution.

Bilberry rhizomes (5) were mostly found at the base of litter horizons (positive side of Axis 1, but not far from the origin) while roots of varying size (7, 8, 9) were mostly found in mineral or organo-mineral horizons (negative side of Axis 1, far from the origin). Dead rhizomes (6) were projected on the negative side of Axis 1, indicating that in the course of time they became buried in the upper part of mineral horizons. Figure 3 shows that in a humus profile at 1870m altitude (micro-podzol) rhizomes were located only in litter horizons, being most abundant in the OH horizon. They were absent from the mineral soil. On the contrary, bilberry roots were increasing in volume from the litter to the mineral soil, being most abundant in the B horizon where ramification occurred, the E horizon being only crossed by vertical roots. Figure 4 shows the distribution of bilberry rhizomes and roots in a humus profile perturbed by rodent activity. Examination of this particular profile revealed that roots colonized preferably mineral horizons, like in the previous case, but also that rhizomes could grow in the mineral soil when loose (backfilling horizon).

Contrary to bilberry, the fine root system of spruce (22, 24, 26, 27, 46) was most abundant in litter, which was confirmed by examination of individual profiles (Figs. 3 and 4). The preference of spruce roots for litter horizons was even still marked than that of bilberry rhizomes. Grass roots (70,

71, 80, 89, 90, 91) were most abundant in mineral horizons (negative side of Axis 1), but in a more shallow position than roots of bilberry, their corresponding points being projected nearer the origin.

Enchytraeid, arthropod and epigeic earthworm faeces (126, 127, 128, 129, 130, 131, 132, 133) were projected on the positive side of Axis 1, approximately at the same level as spruce mycorrhizae, i.e. in OH horizons (Fig. 1). Enchytraeid faeces (126) should be considered as an important component of OH horizons, into which they may constitute up to 24% of the total matrix volume. Earthworm organo-mineral faeces (135, 136) were projected on the negative side of Axis 1, not too far from the origin (Table 1), thus they rather characterized A and "Mound" horizons. Old organo-mineral earthworm faeces (139, 140, 141, 143, 144, 145) were projected roughly at the same level as dead spruce roots (24, 25) and dead bilberry rhizomes (6).

The projection of active and passive variables on Axis 2 indicated a variation in the composition of litter only, the mineral part of the humus profile exhibiting weak variation along this axis. The scaling of the five altitudes along Axis 2 indicated that this axis reflected the effects of elevation on the composition of litter in bilberry carpets. The montane level (950 and 1470m altitude) was characterized by mosses (52, 53, 54, 57, 58, 59), liverworts (55, 56), *Oxalis acetosella* L. (114, 115), *V. vitis-idaea* (111) and *A. alba* (37), while the subalpine level was characterized by *Asteraceae* (102, 105), *L. sylvatica* (62, 63, 64, 65, 66, 67), *Alchemilla alpina* L. (116, 117), *L. decidua* (38, 39) and *P. cembra* (40, 41). *V. myrtillus* (1, 2, 3, 4) and *A. flexuosa* (82, 83, 84, 85, 86, 87, 88) were projected on the negative side of Axis 2, but not far from the origin. Both species can be considered as constant members of bilberry carpets throughout the altitudinal gradient.

The projection of holorganic faecal components typical of OH horizons (126, 127, 128, 130, 131, 132, 133) on the positive side of Axis 2, together with litter components typical of the montane level (37, 52, 53, 54, 55, 56, 57, 58, 59, 111, 114, 115) indicated that OH horizons, typical of moder humus forms (Brêthes et al. 1995), were mostly expressed at low elevation. Thus more organic matter accumulated at the top of soil profiles at the montane compared to the subalpine level. Figures 5 and 6 show that the total thickness of litter horizons and the mean relative volume of animal faeces (only fresh, recognizable material, taken into account) decreased significantly with altitude.

Discussion

Our first work hypothesis was a double strategy for the use of soil nutrients by the subterranean parts of bilberry. The present results, although based on a limited number of samples, pointed to dissimilarities between the vertical distribution of roots, which prevailed in mineral horizons, and rhizomes, which prevailed in organic horizons. We have shown that when the soil was well-aerated, like in the case of rodent mounds, rhizomes were able to penetrate the soil deeply (Fig. 4). This had been also observed in mull humus forms typical of the first (pioneer) stages of colonization by bilberry (André 1994; Bernier & Ponge 1994; Maubon et al. 1995; Ponge et al. 1998). We hypothesize that the main reason for the lesser abundance of bilberry rhizomes in E, B and S horizons and in the A horizon of moder humus is the compact nature of these horizons, due to weak faunal activity, in particular the absence of interconnected pores of enough size (Brêthes et al. 1995). This could explain why the rhizome system of bilberry, the plagiotropism of which has been observed in culture (Barker & Collins 1963, working on the closely related species *Vaccinium angustifolium* Ait., grew worse when it encountered mineral horizons in a mountain slope, resulting in a preferentially downsloping growth (Maubon et al. 1995). Contrary to rhizomes, the diameter of which exceeds one mm near their growing apex (personal observations), the diameter of fine roots of bilberry is less than 0.1mm. Thus the root system of bilberry is better adapted than the rhizome system to penetrate compact mineral horizons where pores created by enchytraeids are of sufficient size (Ponge 1999; Topoliantz et al. 2000).

Our second work hypothesis was that spruce and bilberry were spatially segregated within the soil profile. The present results do not show such segregation, bilberry rhizomes and spruce roots exploiting together the same litter horizons. Nevertheless the root system of bilberry escapes from competition by spruce, by exploiting preferably mineral horizons. Thus in a podzolic soil bilberry will be able to derive nutrients from the illuviation B horizon, where bases leached from litter by colloid transport accumulate (Goldberg et al. 2000), which spruce cannot do due to the scarcity of roots in this horizon (Fig. 3). This point may be of paramount importance if we consider the podzolizing effects of bilberry, probably due to the high production of p-coumaric and protocatechuic acids by senescing

foliage, increasing with altitude (Gallet & Lebreton 1995), to be a mean by which this ericaceous shrub may alleviate competition by spruce and thus may form dense carpets to the detriment of spruce, together with the biochemical inhibition of spruce germination and seedling growth (Gallet 1994; Jäderlund et al. 1996).

The last point to be elucidated was the decrease with altitude of the accumulation of organic matter at the top of soil profiles. This phenomenon, already noted by Bernier (1997), has been confirmed by the present results about litter thickness (Fig. 5) and amount of animal faeces within humus profiles (Fig. 6). Several reasons can be postulated to explain this phenomenon, i) a decrease in primary productivity, ii) a decrease in the recalcitrance of litter to leaching and decomposition, iii) a decrease in the intensity of humification processes. The present study cannot address all these points, since we did not measure plant production nor carbon fluxes in the soil system, but it should be highlighted that we observed a decrease with altitude in the bryophytic component of bilberry carpets, the herb component increasing accordingly. This could be thought to be at the origin of some improvement of litter quality, given the well-known recalcitrance of moss litter towards decomposition (Kilbertus 1968; Ponge 1988), but we must point out that signs of animal activity typical of a rapid disappearance of litter (in particular earthworm activity) did not seem to increase with altitude. In particular all components of faecal material were projected on the positive side of Axis 2 of correspondence analysis (Fig. 1), thus were associated with the montane level rather than with the subalpine level. If we consider that the abundance of animal faeces in humus profiles reflects the level of animal activity (Topoliantz et al. 2000) we must conclude that animal activity decreases with altitude, which indicates a shift from Moder/Mull (both humus forms with a high level of animal activity) to Mor (Ponge et al. 2000; Ponge submitted), despite the observed decrease in litter thickness.

We suspect the leaching of colloidal and soluble organic matter to interact with organic matter accumulation. The decrease in animal activity from the montane to the subalpine level may have far-reaching consequences on humificative processes. If the amount of animal faeces decreases, then more small-sized organic molecules will be leached down the soil profile without being stabilized within big-sized humic assemblages (animal faeces, microbial biomass), according to the podzol model (Kononova 1961; Schoenau & Bettany 1987). Thus a large part of the organic matter produced by

vegetation could escape accumulation at the top of soil profiles while accumulating deeper in B horizons. Given that mosses act as a sponge by retaining water and nutrients coming from precipitation and throughfall (Tamm 1953; Bates 1987), we suspect that organic and mineral molecules present in moss litter cannot be leached so easily than those present in grassy and even bilberry litter (Gallet & Lebreton 1995). Thus both phenomenons, the decrease in the amount of moss litter and the decrease in the deposition of animal faeces may concur synergistically in more leaching of organic matter at the subalpine level, thus in a reduced thickness of accumulated litter.

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Legends of figures

Fig. 1. Correspondence analysis on the data matrix with 159 humus components as active variables and 53 samples as observations. Elevation and depth levels and horizon types were put as additional (passive) variables. Projection of active and passive variables in the plane of the first two factorial axes, extracting 13 and 9% of the total variance, respectively.

Fig. 2. Correlation between Axis 1 coordinates of the 159 humus components and their mean depth (see text and Table 1).

Fig. 3. The influence of profile depth on the abundance of spruce roots and bilberry roots or rhizomes in a micro-podzol (1870m altitude).

Fig. 4. The influence of profile depth on the abundance of spruce roots and bilberry roots or rhizomes in an old rodent mound (1650m altitude).

Fig. 5. Dependence of litter thickness (OL+OF+OH horizons) on altitude in the 12 humus profiles studied.

Fig. 6. Dependence of the mean relative volume of faecal material on altitude in the 12 humus profiles studied.

Table 1. List of humus components found during micromorphological analysis, together with their coordinates along axis 1 of correspondence analysis, their mean vertical distribution (% of soil matrix) and their mean depth

Code	Definition	Axis 1	1cm	2cm	3cm	4cm	5cm	6cm	7cm	8cm	9cm	10cm	11cm	12cm	13cm	14cm	15cm	Mean depth (cm)
1	Entire <i>Vaccinium myrtillus</i> leaf	0.92	2.8	1.9	0.5	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.5
2	Fragmented <i>Vaccinium myrtillus</i> leaf	0.59	1.5	1.5	1.1	0.5	0.4	0.4	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.7
3	<i>Vaccinium myrtillus</i> leaf nerve	0.73	5.5	4.4	2.7	1.8	1.7	1.6	0.5	0.5	0.3	0.2	0.2	0.1	0.1	0.0	0.0	2.6
4	<i>Vaccinium myrtillus</i> stem	0.65	1.7	1.6	0.8	0.7	0.5	0.4	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	2.8
5	Living <i>Vaccinium myrtillus</i> rhizome	0.04	1.3	1.4	1.7	3.3	2.8	2.8	4.1	3.9	3.0	2.6	2.3	1.8	1.6	1.0	1.0	6.4
6	Dead <i>Vaccinium myrtillus</i> rhizome	-0.32	0.1	0.2	0.3	0.4	0.5	0.5	1.4	1.3	0.8	0.6	0.6	0.7	0.7	0.7	0.7	7.3
7	Living <i>Vaccinium myrtillus</i> root < 0.1 mm	-0.45	0.3	0.9	1.5	2.2	2.3	2.5	4.6	5.0	4.4	4.1	4.4	3.6	3.4	2.6	2.6	8.6
8	Living <i>Vaccinium myrtillus</i> root 0.1-0.3 mm	-0.51	0.3	0.5	0.8	1.1	1.0	1.2	1.7	1.6	1.8	1.8	1.8	1.9	1.9	1.8	1.8	8.8
9	Living <i>Vaccinium myrtillus</i> root > 0.3 mm	-0.88	0.1	0.0	0.2	0.4	0.4	0.5	0.7	0.7	0.7	1.0	1.1	1.0	0.9	1.3	1.3	9.3
10	Dead <i>Vaccinium myrtillus</i> root < 0.1 mm	-0.63	0.1	0.2	0.7	1.2	1.5	1.7	2.5	2.4	2.3	2.3	2.4	2.4	2.3	3.2	3.2	9.3
11	Dead <i>Vaccinium myrtillus</i> root 0.1-0.3 mm	-0.72	0.0	0.0	0.0	0.1	0.2	0.3	0.4	0.4	0.4	0.4	0.5	0.4	0.5	0.6	0.6	10.0
12	Dead <i>Vaccinium myrtillus</i> root > 0.3 mm	-0.86	0.0	0.0	0.0	0.1	0.1	0.1	0.2	0.2	0.2	0.2	0.3	0.3	0.3	0.3	0.3	9.7
13	Fragment of <i>Vaccinium myrtillus</i> root or rhizome	-0.03	0.3	0.4	0.5	0.4	0.2	0.2	0.3	0.3	0.3	0.2	0.2	0.2	0.2	0.2	0.2	7.1
14	Intact <i>Picea</i> needle	0.93	7.9	7.4	3.0	2.3	1.8	0.4	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	1.8
15	Fragmented or degraded <i>Picea</i> needle	0.73	2.5	2.6	3.4	2.9	2.2	1.9	0.7	0.5	0.2	0.1	0.1	0.1	0.0	0.0	0.0	3.3
16	Coniferous bark	0.24	1.6	1.7	2.4	2.5	2.3	2.1	2.0	2.7	2.7	2.7	2.6	1.5	0.7	0.6	0.6	5.4
17	Coniferous wood	0.00	0.7	0.7	0.6	0.7	0.5	1.7	1.7	1.6	1.6	1.6	0.1	0.1	0.1	0.0	0.0	4.2
18	Charcoal	0.00	0.1	0.1	0.1	0.1	0.2	0.2	0.4	0.4	0.4	0.2	0.2	0.1	0.1	0.2	0.2	9.0
19	Coniferous pollen	0.74	0.7	0.5	0.4	0.4	0.4	0.4	0.2	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	3.6
20	Scale from coniferous male cone	0.97	2.8	2.6	0.4	1.1	1.1	1.0	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	2.5
21	Coniferous twig	0.83	1.1	1.1	0.8	0.7	0.7	0.5	0.4	0.4	0.4	0.3	0.1	0.0	0.0	0.0	0.0	2.8
22	Living <i>Picea</i> root < 0.5 mm	0.09	0.1	0.2	0.5	0.4	0.5	0.5	0.8	0.9	0.7	0.4	0.4	0.5	0.4	0.2	0.2	6.9
23	Living <i>Picea</i> root > 0.5 mm	-0.14	0.2	0.2	0.5	0.6	0.7	0.8	1.2	1.7	1.2	1.2	1.1	1.1	0.6	0.5	0.5	6.7
24	Dead <i>Picea</i> root < 0.5 mm	-0.26	0.1	0.2	0.3	0.4	0.6	0.7	1.1	1.1	1.1	0.9	0.9	0.9	0.6	0.4	0.4	7.2
25	Dead <i>Picea</i> root > 0.5 mm	-0.16	0.1	0.2	0.4	0.4	0.7	0.8	1.2	1.2	1.1	0.9	0.7	0.6	0.4	0.3	0.3	7.1
26	Living <i>Picea</i> mycorrhiza	0.45	0.7	1.2	2.0	1.9	2.0	1.7	1.3	1.1	0.8	0.4	0.3	0.4	0.3	0.2	0.2	5.6
27	Dead <i>Picea</i> mycorrhiza	0.10	0.3	0.5	1.6	1.7	3.0	2.9	4.0	3.7	3.7	2.9	2.3	2.3	1.0	0.1	0.1	7.0
28	Fragment of coniferous root	0.24	0.1	0.1	0.2	0.5	0.2	0.2	1.1	1.1	1.0	0.1	0.1	0.1	0.2	0.2	0.2	8.0
29	Living <i>Abies</i> root < 0.5 mm	0.82	0.1	0.1	0.1	0.2	0.2	0.5	0.5	0.5	0.5	0.5	0.3	0.0	0.0	0.0	0.0	5.5
30	Living <i>Abies</i> root > 0.5 mm	0.66	0.1	0.1	0.1	0.1	0.1	0.7	0.7	0.7	0.7	0.7	0.7	0.0	0.0	0.0	0.0	4.4
31	Living <i>Larix</i> root < 0.5 mm	-0.71	0.0	0.0	0.2	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.1	0.1	0.1	0.1	7.0
32	Living <i>Larix</i> root > 0.5 mm	-0.60	0.1	0.1	0.2	0.5	0.5	0.5	0.5	0.5	0.4	0.4	0.4	0.1	0.1	0.1	0.1	6.8
33	Dead <i>Larix</i> root < 0.5 mm	0.52	0.0	0.0	0.3	0.5	0.4	0.7	0.7	0.7	0.7	0.6	0.1	0.1	0.1	0.1	0.1	7.8
34	Dead <i>Larix</i> root > 0.5 mm	0.16	0.1	0.1	0.2	0.2	0.3	0.4	0.4	0.4	0.4	0.4	0.2	0.2	0.2	0.2	0.2	7.0
35	Dead <i>Larix</i> mycorrhiza	0.75	0.7	0.7	0.5	0.1	0.2	0.2	0.2	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	5.5
36	Living <i>Larix</i> mycorrhiza	0.76	1.5	1.6	0.4	0.5	0.4	0.5	0.5	0.5	0.4	0.4	0.3	0.0	0.0	0.0	0.0	3.9
37	Entire <i>Abies</i> needle	1.05	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.5
38	Entire <i>Larix</i> needle	0.80	3.4	1.3	0.6	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.4
39	Fragmented <i>Larix</i> needle	0.56	0.5	1.5	1.0	0.3	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.7
40	Entire <i>Pinus</i> needle	0.75	0.4	0.4	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.5
41	Fragmented <i>Pinus</i> needle	0.45	0.1	0.1	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.4
42	Living <i>Pinus</i> mycorrhiza	-0.01	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.4
43	Dead <i>Pinus</i> mycorrhiza	0.25	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.0
44	Living <i>Pinus</i> root	0.70	0.1	0.1	0.1	0.1	0.0	0.2	0.2	0.2	0.2	0.2	0.2	0.0	0.0	0.0	0.0	4.6
45	Dead <i>Pinus</i> root	0.81	0.5	0.5	0.7	0.6	0.5	0.5	0.5	0.5	0.5	0.5	0.4	0.0	0.0	0.0	0.0	3.8
46	<i>Cenococcum geophilum</i>	0.64	0.7	0.9	1.2	1.1	1.1	0.7	0.6	0.6	0.4	0.2	0.2	0.3	0.2	0.0	0.0	4.7
47	White rhizomorph	0.70	1.7	1.8	1.6	1.3	1.1	1.3	1.0	0.9	0.5	0.5	0.5	0.3	0.2	0.0	0.0	4.2
48	Brown rhizomorph	0.64	0.8	0.8	0.5	0.6	0.6	0.5	0.5	0.4	0.2	0.2	0.1	0.1	0.1	0.0	0.0	4.8
49	Hyaline mycelium	0.28	0.5	0.6	0.8	0.8	0.8	0.8	0.7	0.6	0.3	0.3	0.3	0.3	0.4	0.3	0.3	6.0
50	Dematiaceae mycelium	0.48	0.9	0.8	1.0	1.0	0.8	0.8	0.5	0.4	0.1	0.2	0.1	0.3	0.2	0.1	0.1	5.1
51	<i>Marasmius androsaceus</i>	0.63	0.5	0.5	0.3	0.2	0.1	0.1	0.2	0.2	0.2	0.1	0.1	0.1	0.0	0.0	0.0	5.3
52	Stemlet of <i>Hylocomium splendens</i>	0.96	5.5	4.7	2.2	1.3	1.1	1.1	0.5	0.5	0.4	0.4	0.1	0.0	0.0	0.0	0.0	2.5
53	Stemlet of <i>Rhytidiadelphus triqueter</i>	1.05	9.4	9.3	7.0	4.4	4.2	4.2	2.8	2.8	2.7	2.7	0.0	0.0	0.0	0.0	0.0	2.8
54	Fragmented moss leaf	0.84	3.3	2.9	1.6	0.6	0.3	0.3	0.2	0.2	0.2	0.1	0.1	0.4	0.3	0.0	0.0	3.3
55	Fragment of liverwort	0.70	0.8	1.4	3.1	3.2	3.1	1.5	1.5	1.5	1.5	1.5	0.7	0.0	0.0	0.0	0.0	3.3
56	Liverwort rhizoid	1.13	2.3	2.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.6
57	Fragmented bryophytic rachis	0.68	5.9	5.4	5.4	3.6	2.7	2.7	2.2	1.7	0.7	0.3	0.3	1.0	1.0	0.2	0.2	4.3
58	Leafed stalk of <i>Hylocomium splendens</i>	1.03	1.8	1.2	0.5	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.6
59	Leafed stalk of <i>Rhytidiadelphus triqueter</i>	1.11	1.7	1.1	0.7	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.5
60	Lichen	0.84	0.7	0.7	0.2	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.4
61	Living <i>Luzula</i> leaf	0.82	0.5	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.5
62	Dead <i>Luzula</i> leaf	0.71	2.4	0.7	0.3	0.3	0.3	0.3	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.7
63	Fragmented <i>Luzula</i> leaf	0.21	2.0	1.3	2.4	1.8	1.9	1.9	1.2	1.2	0.4	0.4	0.4	0.2	0.2	0.2	0.2	4.7
64	Dead <i>Luzula</i> stem	0.75	0.2	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.8
65	Living <i>Luzula</i> stem	0.82	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.0
66	Living <i>Luzula</i> foliar base	-0.18	0.5	0.7	0.0	0.0	0.1	0.1	0.0	0.0	0.3	0.3	0.3	0.3	0.3	0.3	0.3	4.2
67	Dead <i>Luzula</i> foliar base	0.17	0.5	0.1	0.7	0.7	0.6	0.6	0.6	0.6	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.6
68	Living <i>Luzula</i> rhizome	0.22	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6
69	Dead <i>Luzula</i> rhizome	0.24	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.7
70	Living <i>Luzula</i> root < 0.3 mm	-0.17	0.8	1.2	2.0	2.2	2.2	1.9	1.9	1.9	0.9	0.9	0.9	0.6	0.7	0.6	0.6	6.1
71	Living <i>Luzula</i> root > 0.3 mm	-0.07	0.1	0.2	0.3	0.4	0.3	0.3	0.4	0.4	0.2	0.2	0.2	0.2	0.1	0.1	0.1	7.9
72	Dead <i>Luzula</i> root < 0.3 mm	-0.43	0.2	0.4	1.3	1.1	1.1	1.1	1.2	1.2	0.7	0.7	0.7	0.5	0.5	0.5	0.5	6.4
73	Dead <i>Luzula</i> root > 0.3 mm	-0.48	0.1	0.1	0.1	0.1	0.1	0.1	0.3	0.3	0.2							

Table 1. (continued)

Code	Definition	Axis 1	1cm	2cm	3cm	4cm	5cm	6cm	7cm	8cm	9cm	10cm	11cm	12cm	13cm	14cm	15cm	Mean depth (cm)
101	<i>Homogyne</i> flower stem	0.74	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.4
102	Fragmented <i>Asteraceae</i> leaf	0.43	0.4	0.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.3
103	Living <i>Asteraceae</i> root	-0.11	0.1	0.4	0.3	0.3	0.3	0.3	0.3	0.3	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.6
104	Dead <i>Asteraceae</i> root	-0.17	0.0	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	6.1
105	Living <i>Asteraceae</i> leaf base	0.15	0.2	0.6	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.8
106	<i>Vaccinium vitis-idaea</i> leaf	0.42	0.0	0.1	0.1	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.3
107	Living <i>Vaccinium vitis-idaea</i> root	-0.03	0.1	0.1	0.1	0.2	0.1	0.1	0.2	0.2	0.2	0.2	0.2	0.2	0.1	0.0	0.0	7.0
108	Dead <i>Vaccinium vitis-idaea</i> root	0.03	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	6.8
109	Living <i>Vaccinium vitis-idaea</i> rhizome	0.55	0.0	0.0	0.2	0.3	0.0	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	5.1
110	Dead <i>Vaccinium vitis-idaea</i> rhizome	0.82	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.1
111	Fragmented <i>Vaccinium vitis-idaea</i> stem	0.47	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.3
112	Living <i>Oxalis</i> root	0.82	0.2	0.2	0.2	0.4	0.4	0.4	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	3.9
113	Dead <i>Oxalis</i> root	0.96	0.2	0.2	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	4.0
114	Living <i>Oxalis</i> leaf base	0.90	0.1	0.0	0.1	0.2	0.2	0.2	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.1
115	Dead <i>Oxalis</i> leaf base	0.91	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.5
116	Fragmented <i>Alchemilla</i> leaf base	0.36	0.0	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.6
117	<i>Alchemilla</i> miscellaneous fragment	0.02	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.2
118	Living <i>Alchemilla</i> root	0.14	0.0	0.1	0.0	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.2
119	Dead <i>Alchemilla</i> root	-0.69	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	8.5
120	Herbaceous plant tissue	0.21	0.4	1.0	0.9	0.9	1.1	1.1	0.6	0.6	0.2	0.2	0.2	0.2	0.1	0.2	0.2	6.5
121	Spore	0.29	0.1	0.2	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	4.8
122	Bud fragment	0.70	0.2	0.2	0.2	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	2.8
123	Exuvia	0.70	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.5
124	Miscellaneous plant fragment	0.46	0.0	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.8
125	Animal hair	0.78	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	3.3
126	Organic enchytraeid faeces	0.45	1.9	3.4	6.8	10.1	7.8	7.9	6.1	5.6	4.3	3.0	2.2	2.0	1.9	0.6	0.6	5.2
127	Organic epigeic earthworm faeces	0.64	2.3	3.1	5.0	4.2	4.3	3.2	2.0	1.8	1.2	1.0	0.4	0.2	0.2	0.1	0.1	4.1
128	Organic silver faecal material	0.56	1.0	1.2	2.6	2.7	3.7	3.7	2.0	1.3	1.2	1.0	0.7	0.4	0.3	0.2	0.2	4.8
129	Organic collembolan faeces	0.55	0.2	0.3	0.3	0.2	0.2	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	3.6
130	Organic plate-like faeces	0.61	0.4	0.4	0.3	0.3	0.2	0.1	0.1	0.1	0.2	0.2	0.2	0.1	0.1	0.2	0.2	5.4
131	Organic insect larvae faeces	0.36	0.0	0.0	0.2	0.2	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.1	0.1	5.2
132	Organic mite faeces	0.48	0.1	0.1	0.1	0.4	0.0	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.7
133	Organic millipede faeces	0.67	1.0	1.0	1.2	0.9	0.8	0.6	0.4	0.2	0.2	0.2	0.1	0.0	0.0	0.2	0.2	3.4
134	Organo-mineral enchytraeid faeces	0.20	0.2	0.2	3.6	3.9	4.3	4.6	4.6	4.8	4.3	4.3	2.7	1.2	1.0	1.0	1.0	7.6
135	Organo-mineral endogeic earthworm faeces	-0.58	0.1	0.9	1.5	2.2	2.4	2.7	2.8	3.1	2.5	2.5	2.8	2.4	2.2	2.2	2.2	8.3
136	Organo-mineral anecic earthworm faeces	-0.38	0.0	0.0	0.6	1.4	1.4	1.3	1.3	1.3	1.3	1.3	1.3	0.6	0.5	0.4	0.4	6.6
137	Mineral endogeic earthworm faeces	-0.76	0.0	0.1	0.8	0.9	0.9	1.7	2.0	2.8	2.7	2.7	2.7	2.0	1.9	3.3	3.3	8.9
138	Mineral enchytraeid faeces	-0.79	0.0	0.1	0.8	1.0	1.0	1.0	1.7	2.1	3.4	3.4	3.3	2.6	2.0	1.6	1.6	8.2
139	Organo-mineral granular aggregates	-1.36	0.0	0.0	0.2	0.3	0.5	0.5	0.5	0.5	0.8	0.8	1.1	1.2	1.8	1.4	1.4	11.3
140	Organo-mineral crumb aggregates	-1.15	0.1	0.1	0.2	0.4	0.4	0.4	0.4	0.4	0.9	1.0	1.1	1.4	1.9	3.3	3.3	11.8
141	Organo-mineral polyedric aggregates	-1.27	0.0	0.0	0.0	0.6	0.7	0.7	1.2	1.2	1.4	1.4	2.5	1.9	1.8	2.3	2.3	10.5
142	Mineral particulate aggregates	-1.70	0.0	0.0	0.0	0.0	3.8	3.8	5.7	5.7	7.5	9.4	9.4	9.4	8.1	4.7	4.7	11.7
143	Mineral granular aggregates	-1.60	0.0	0.0	0.0	0.2	0.3	0.3	0.5	0.5	0.6	1.8	1.8	2.0	1.8	0.8	0.8	11.7
144	Mineral crumb aggregates	-1.41	0.0	0.0	0.1	0.4	0.4	1.0	1.0	1.0	1.3	3.0	3.8	4.5	6.0	5.4	5.4	12.7
145	Mineral polyedric aggregates	-1.51	0.0	0.0	0.0	0.0	0.1	0.1	0.1	0.1	0.9	1.5	1.8	1.8	2.9	2.9	2.9	12.1
146	Coarse gravel	-1.53	0.0	0.0	0.0	0.2	0.6	0.6	0.9	0.9	2.2	2.3	2.4	2.8	3.4	5.2	5.2	12.2
147	Fine gravel	-1.48	0.0	0.0	0.0	0.1	0.2	0.2	0.7	0.7	1.9	2.0	2.1	2.3	2.2	2.5	2.5	11.7
148	Coarse sand	-1.42	0.0	0.1	0.1	0.4	0.4	0.4	0.6	0.6	1.2	1.4	1.4	1.7	2.1	4.0	4.0	12.3
149	Fine sand	-1.28	0.0	0.1	0.2	0.6	0.6	0.6	0.8	0.8	1.6	1.6	1.7	2.3	2.3	2.1	2.1	11.1
150	Silt	-1.37	0.0	0.0	0.5	0.9	0.9	0.9	1.8	1.8	3.2	3.2	3.4	3.5	3.6	3.2	3.2	11.0
151	Boulder	-1.49	0.0	0.0	0.3	1.4	1.7	1.7	2.9	2.9	5.8	6.4	6.5	6.6	9.8	13.6	13.6	11.8
152	Quartz grain	-0.81	0.1	0.1	0.3	0.2	0.2	0.3	0.3	0.5	0.9	1.0	0.9	0.8	0.8	1.2	1.2	7.8
153	Mineral fragment	-0.66	0.0	0.0	0.1	0.1	0.1	0.0	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.2	0.2	7.3
154	Enchytraeid	0.05	0.0	0.0	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	6.6
155	Epigeic earthworm	0.64	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.6
156	Collembolan	0.71	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	4.5
157	Mite	0.81	0.1	0.1	0.1	0.1	0.1	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	2.8
158	Larva	-0.84	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	6.3
159	Millipede	0.86	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	1.5

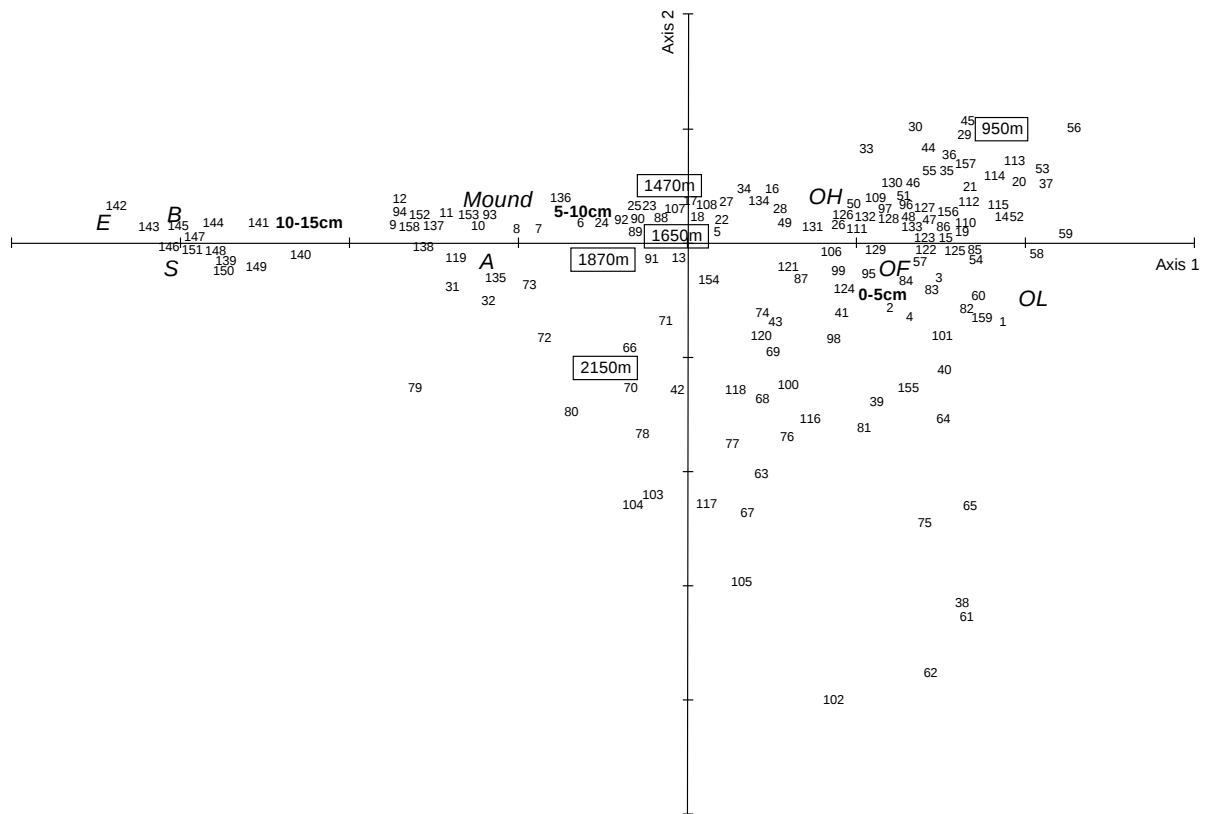


Fig. 1

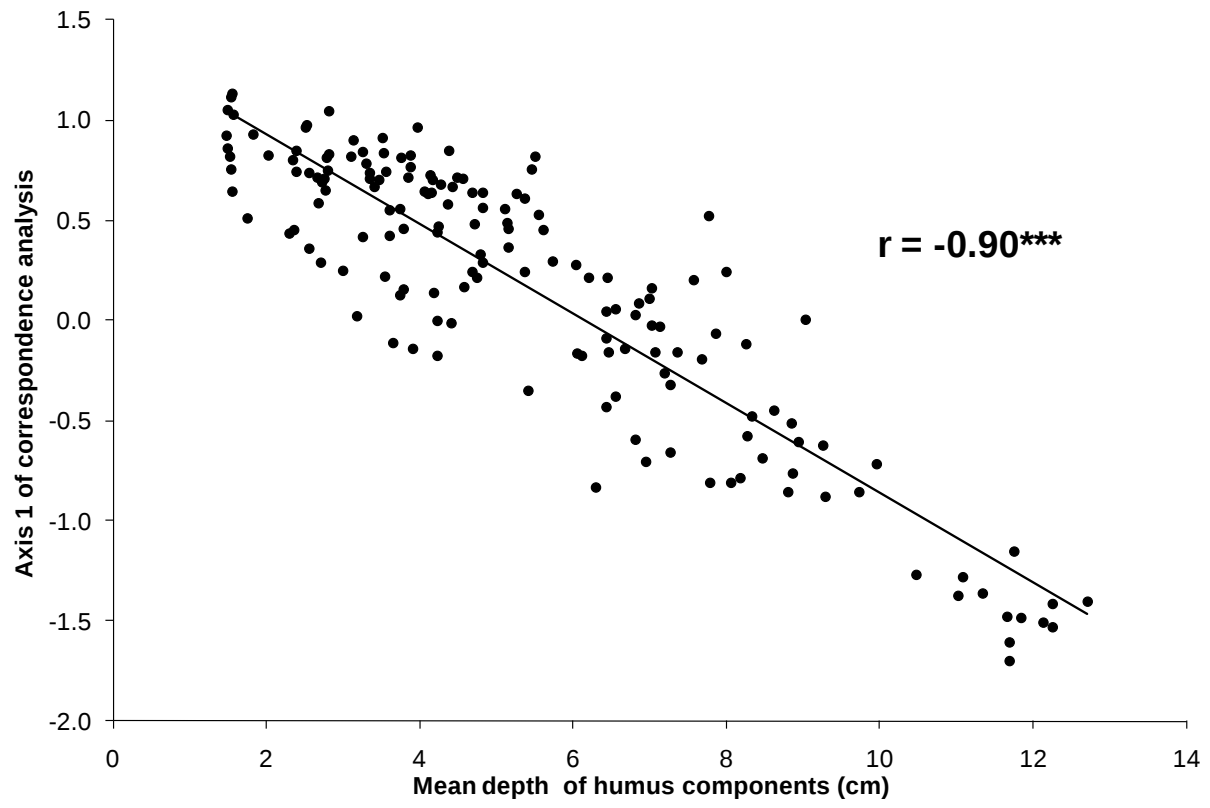


Fig. 2

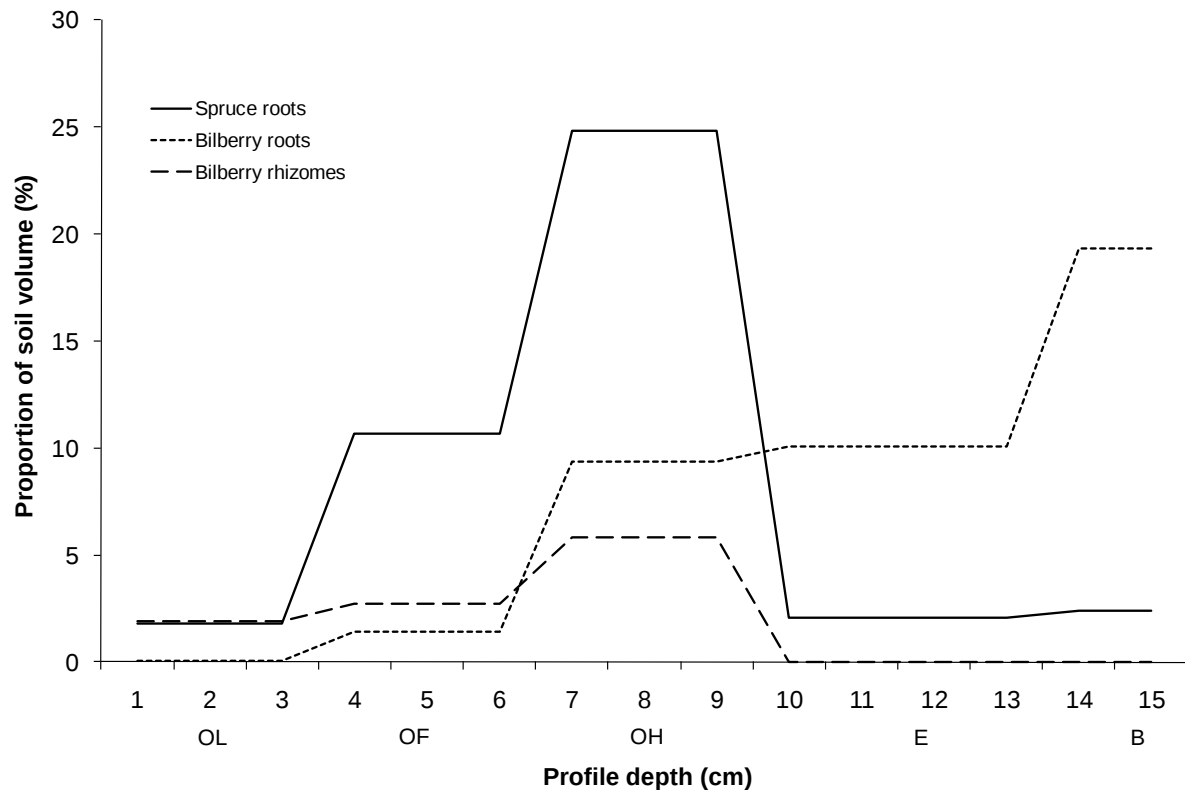


Fig. 3

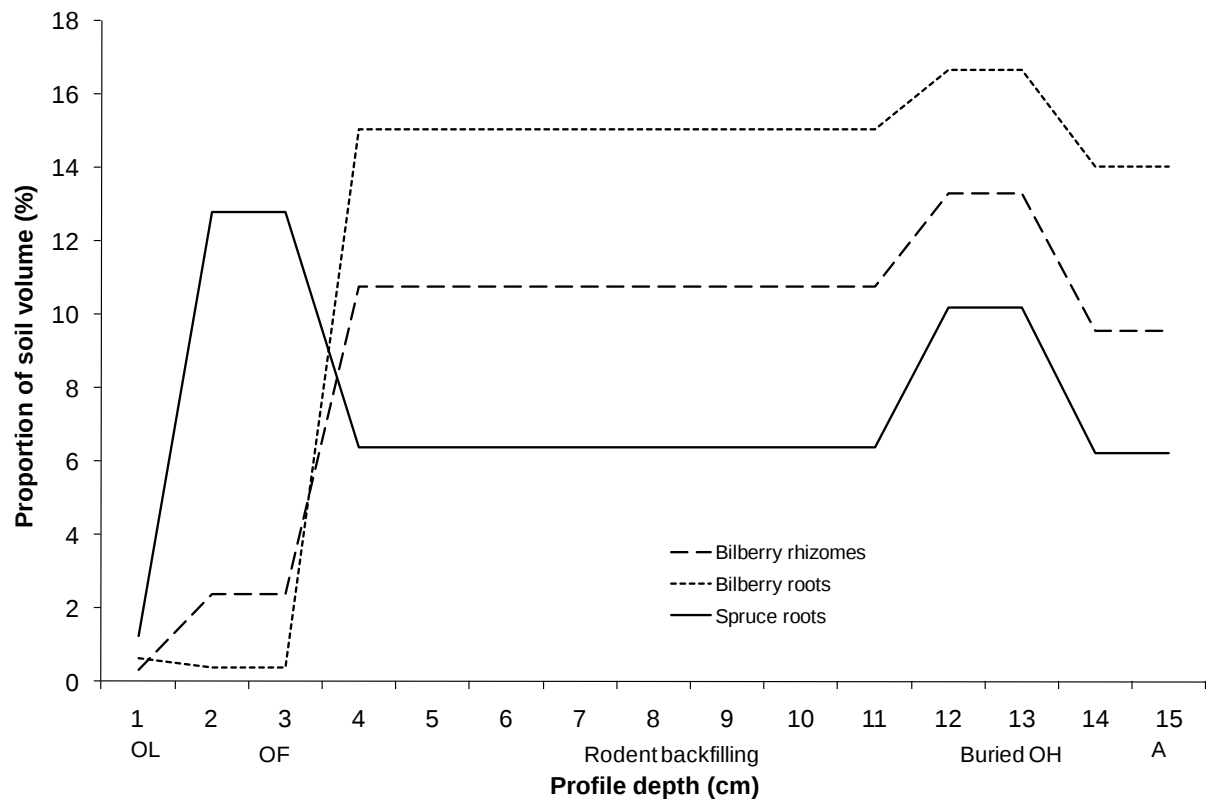


Fig. 4

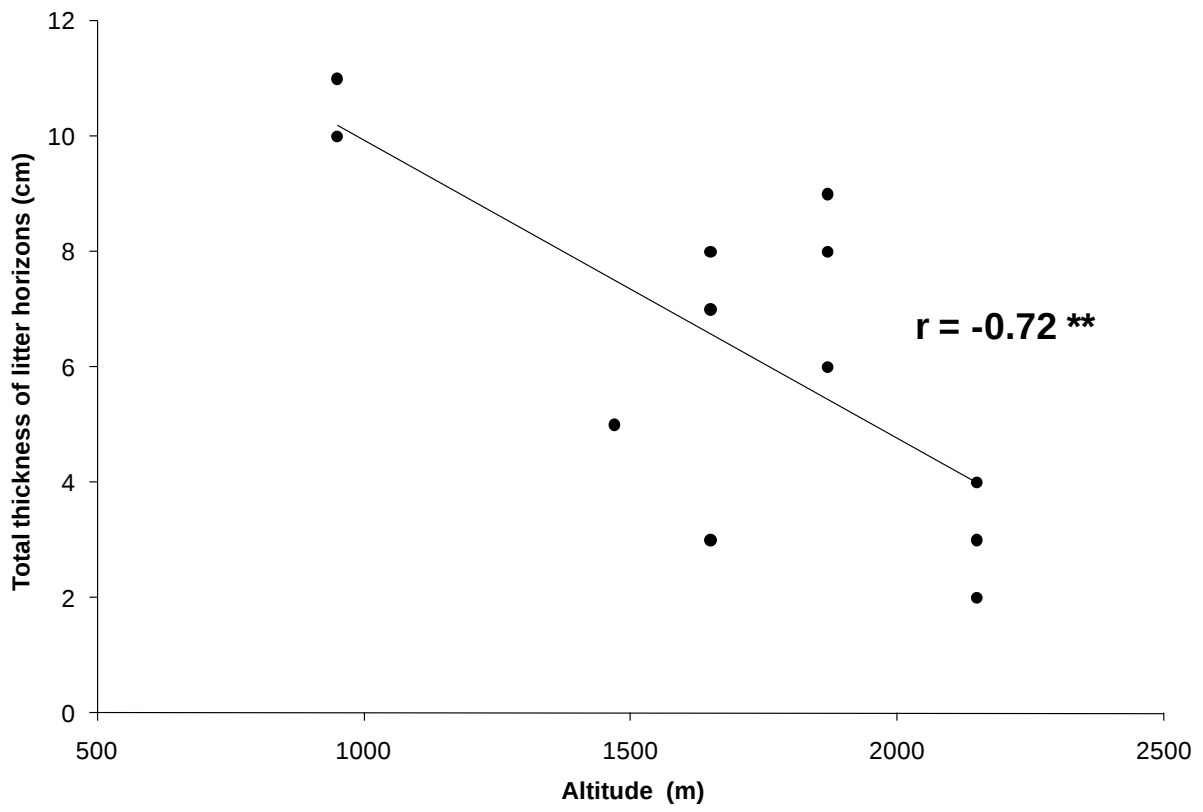


Fig. 5

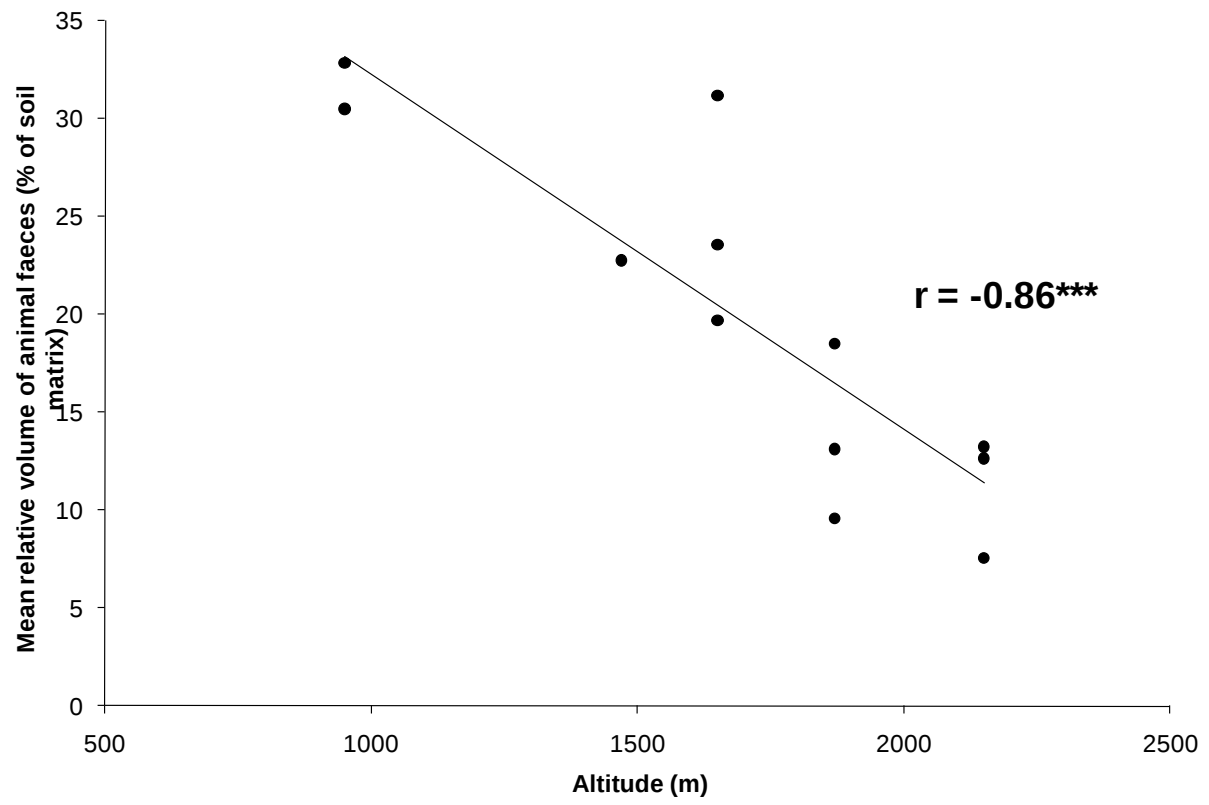


Fig. 6