Tree species identity influences the vertical distribution of labile and recalcitrant carbon in a temperate deciduous forest soil.

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Abstract

In terrestrial environments, soil organic matter (SOM) is the largest organic carbon (C) pool. The quantity and quality of organic carbon in soils can be affected by vegetation through influencing the inputs and outputs of SOM. We examined how storage and quality of C in SOM were affected by vegetation under grass cover or single and a polyculture plot of Betula pendula, Alnus glutinosa and Fagus sylvatica. An acid hydrolysis approach was used to quantify three SOM fractions differing in biodegradability. Tree species identity and stand composition had no significant effect on the total amount of C stored in different SOM fractions to a depth of one meter. However, when examining individual SOM fractions in the upper layers of the soil profile, significantly more C was stored in the putatively more labile fractions 1 and 2 under F. sylvatica and A. glutinosa, respectively. In deeper soil layers, the highest storage of recalcitrant organic C was found under the tree polyculture. The vertical distribution of these three soil organic C pools was compared to C inputs via decomposed leaf litter. Our data indicated that in the tree species polyculture, combining litter inputs of multiple species can have a positive impact on the accumulation of acid resistant recalcitrant C in deep soil layers in 4 years. This C fraction has the greatest potential for long-term sequestration.

Keywords Fractionation; Acid hydrolysis; Polyculture; Carbon storage; Tree species mixture; Decomposition
1. Introduction

Soil organic matter (SOM) represents the largest reservoir of terrestrial organic carbon (C) on Earth, and the organic residues that comprise SOM range from relatively intact plant or microbial material to highly decomposed humic substances (Rumpel et al., 2002). Through differences in litter quality, plant species have potential to influence the storage and dynamics of C in soils, as reported in several previous studies (Binkley and Valentine, 1991; Hagen-Thorn et al., 2004; Jandl et al., 2007; Leuchner et al., 2013). However, there is growing evidence that molecular structure is less important than previously believed as a factor controlling the formation of SOM, and that the inputs from roots and microbial degradation products are more important than previously assumed (Schmidt et al., 2011). However, it remains undisputed that these factors result in a heterogeneous mixture of organic compounds (von Lützow et al., 2006; Schmidt et al., 2011; Tfaily et al., 2015).

To investigate the composition of soil organic matter, SOM can be fractionated into several pools using a range of techniques (e.g. Paul et al., 2001; Weil et al., 2003; Gregorich et al., 2003; Bajgai et al., 2013). Each of these methods attempts to isolate soil organic matter pools of different longevity. A common method used, is to separate pools on the basis of biodegradability (Rovira and Vallejo, 2007). This method separates soil organic matter based on acid solubility into labile and recalcitrant pools, which are believed to have different turnover times (McLauchlan et al., 2004). For example, labile C fraction composed of compounds such as soluble sugars, starch, and other carbohydrates, has been show to play a dominant role in the evolution of CO₂ from soil due to preferential decomposition and rapid turnover (Belay-Tedla et al., 2009). In contrast the recalcitrant fraction is thought to degrade slowly, thus contributing to long term C storage in soils. For
example, lignified humus and some physically protected labile SOM can be retained in soils for several thousand years (Zou et al., 2005; Dungait et al., 2012; Kellner et al., 2014). However, how the composition of aboveground vegetation affects the distribution of these pools in soil systems is largely unknown (De Deyn et al., 2008).

Tree species may influence soil organic carbon (SOC) stocks through a variety of mechanisms such as: (i) differences in net primary productivity and the production of detritus (Montagnini et al., 1993; Hansen et al., 2009), (ii) variation in the quality and complexity of organic matter input to soils detritus originating from leaf, root and mycorrhizal biomass (Hagen-Thorn et al., 2004), (iii) variation in the depth and distribution of roots (Carvalheiro and Nepstad, 1996; Lai et al., 2015), and (iv) by altering soil invertebrate and microbial populations (Hobbie et al., 2006; Lynch et al., 2012). Soil organic carbon accumulation may differ in species-diverse communities, compared to monocultures through variation in biomass inputs to soils, and SOM transformation processes in the soil (Steinbeiss et al., 2008). In mixed species communities, species interactions may either increase productivity through resource use complementarity (Loreau and Hector, 2001; Richards and Schmidt, 2010) or decrease productivity through inter- or intra-specific competition (Carnus et al., 2006). Plant community composition in mixtures may also influence long term storage of soil C through species-specific differences in plant detritus chemical composition and input rates (Six et al., 2002). In addition, interaction between litter types in mixed species communities may affect rates of decomposition and turnover (King et al., 2002). This is particularly true in natural forest ecosystems, where the litter layer can be comprised of inputs from many species, thus forming more complex organic substances compared to the litter layer of mono-specific forests. The highly complex and heterogeneous organic residues found in the SOM of mixed
communities have been shown to alter soil C residence times through differences in biodegradability (Sollins et al., 1996).

Our study determined the vertical distribution of labile and recalcitrant C fractions in SOM that occurred within the initial four years of forest establishment. We examined the effects of vegetation on the storage of C fractions in soils under grass or trees of A. glutinosa, B. pendula, F. sylvatica grown in monoculture, or a polyculture of the three tree species. B. pendula is a light-demanding, early successional species with fast juvenile growth (Fischer et al., 2002). A. glutinosa is an N-fixing, water-demanding pioneer species, also with high juvenile growth rates (Braun, 1974). Lastly, F. sylvatica is shade tolerant and slow growing when juvenile (Ellenberg et al., 1991), can persist in the understory, and often dominates late successional forest. We hypothesized that in the long term SOC storage would be positively affected by growing trees species selected for contrasting functional traits in polyculture.

2. Materials and Methods

2.1 Study area

The BangorDIVERSE experimental site was established at Henfaes Research Centre, Bangor University, North Wales, UK (53°14′ N, 4°01′W) in March 2004 on two fields with a total area of 2.36 ha. Soils are fine loamy brown earth over gravel (Rheidol series) and classified as Fluventic Dystrochrept in the USDA system (Smith et al., 2013a). Soil texture in the 0-10 cm layer was 48.2 ± 1.3 % sand, 33.6 ± 0.9 % silt and 18.2 ± 2.1 % clay, determined by laser diffraction (Coulter LS particle size analyser). The soil pH is 5.4 in the 0-10 cm layer increases to 6.3 at 100 cm soil depth. Soil physicochemical properties are shown in Suppl. Table 1. Climate at the
site is classified as hyperoceanic. Mean annual temperature throughout 2005–2008 was 11.5 °C with an annual rainfall of 1034 mm (Campbell Scientific Ltd, Shepshed, UK).

2.2 Plantation design

Before tree planting, all vegetation was removed from the fields including the grass plots and the soil was ploughed and raked. Plots were established in four replicated blocks of single species or two and three species mixtures of Alnus glutinosa L., Betula pendula Roth., Fagus sylvatica L., Fraxinus excelsior L., Acer pseudoplatanus L., Castanea sativa Mill. and Quercus robur L. The trees were selected due to their contrasting shade tolerance, successional chronology and to represent a range of taxonomic, physiological and ecological types. Two blocks were sited in each field and the minimum distance between any two plots of the same composition was 35 m.

The size of the plots were 62 m² for the grassland, 81 m² for single species, and 121 and 196 m² for the two species and three species plots respectively. A replacement series design (with inter-tree spacing constant between treatments) was selected because of the experiments objective of being realistic in reflecting the practical realities of how forests comprising monocultures or mixtures of potential canopy tree species could be established (Jolliffe, 2000). The site was planted with 60 cm saplings of each species with an inter-tree spacing of 1 m (10,000 stems ha⁻¹). A systematic hexagonal planting design (Aguiar et al., 2001) was used to maximise the mixing effect so that, in the three species mixture blocks, each tree was surrounded by nearest neighbours of two-con-specific individuals and one and three individuals of the other two species, respectively, resulting in each tree having six equidistant neighbours. On the grassland plots, a grass cover was allowed to regenerate from remnants to form a
sward composed of a mixture of *Lolium perenne* L., *Dactylis glomerata* L. and *Agrostis stolonifera* L. In the work reported here, we used the plots of *A. glutinosa*, *B. pendula* and *F. sylvatica* and a three species polyculture of these species.

2.3 Positioning of the plots used

The initial soil organic matter was determined in the top 0-10 cm layer on a 10 x 10 m grid. The mean SOM content across the fields was ca. 6 %, but across the fields varied between 4 to 8 % (Fig. 1). Historically, both fields were pasture, but since the 1980s one field (field 1) was used for small scale forestry experiments, while in 2003 the other field (field 2) was ploughed and planted with oil seed rape. Two blocks each containing a replicate single species and mixed species plot as well as the associated grassland plot, were located on the field used in 2003 for soil seed rape. In the other field one block was positioned on an area previously planted with *Salix* as short rotation coppice trial, while the other block was positioned on an area previously used to grow mainly *Q. robur* and *F. sylvatica* saplings. In 2008, one of the *A. glutinosa* plots was damaged. To balance the number of replicates, one plot each of the *F. sylvatica*, *B. pendula* and polyculture were removed from the analysis by random selection, leaving the distribution of plots used as shown in Figure 1.

2.4 Soil collection and sample processing

Soil was collected by excavating 100 x 100 x 100 cm pits in the centre of each grassland plot and each of the three tree species monoculture and tree three species polyculture plots in September of 2008 (15 pits in total). In the polyculture plots, samples were collected from a pit at an equal distance from *A. glutinosa*, *B. pendula* and *F. sylvatica*. Soil samples were collected from seven layers (0-10, 10-20, 20-30,
A subset of four layers (0-10, 10-20, 40-50 and 90-100 cm) were then used for C fractionation using acid hydrolysis. To ensure representativeness, samples were obtained from each layer, approximately 100 g of soil was collected from each side of the soil pit and mixed to produce a composite 400 g sample. Soils were then air dried, carefully homogenized and sieved to pass through a 2 mm sieve, a 50 g sub-sample was then taken and finely ground using a ball mill (Retsch Mixer Mill MM 200) and passed through a 100 µm sieve prior to acid hydrolysis. Soil pH and electrical conductivity were determined in a 1:2 v/v slurry of soil and distilled water with standard electrodes. Moisture content was determined after drying at 105 °C for 72 h, and organic matter as loss-on-ignition at 450 °C for 16 h. Bulk density was determined using 100 cm$^3$ cores and corrected for stone content (Rowell, 1994). The clay content of soils was determined by a simplified method combining wet sieving and sedimentation steps, as proposed by Kettler et al., (2001). Soil microbial biomass C of surface soil (0-10 cm) was determined according to the CHCl$_3$ fumigation-extraction method of Vance et al., (1987).

### 2.5 Leaf and root sample collection

Fully expanded leaves exposed to full incident light, were collected from the outside of the crown at upper and middle of the canopy positions. Five trees of each species were randomly selected and approximately 2 g of leaves were collected and combined into a composite sample (ca. 10 g) for each species. Root samples were collected using an 8 cm diameter soil corer. Grassland sward was collected from three undisturbed locations, washed and the leaves and roots separated before being oven dried at 80 °C for 24 h. Dried leaves and roots were separately ground using a ball mill (Retsch Mixer Mill MM 200) and passed through a 100 µm sieve. The total C
content of soil and plant materials was determined using a TruSpec® CN analyser (Leco Corp., St Joseph, MI). The leaf and root materials were also analysed by sequential acids hydrolysis as described below.

2.6 Acid hydrolysis of soil and plant materials

Chemical methods allow fractionation of SOC into pools of putative identity. For example in a two-step acid hydrolysis approach, the labile fraction is further divided into two pools- labile fraction 1 that could be putatively identified as polysaccharides, derived from plant and microbial sources, and labile fraction 2 that contains cellulose which is more resistant than fraction 1 (Rovira and Vallejo, 2007). Labile soil C (fraction 1) was extracted from 0.5 g of air dried soil, which was taken from the homogenised 50 g sample described above. The soil was transferred into a sealable Pyrex tube and 15 ml of 2.5 M H₂SO₄ was added and thoroughly mixed. The mixture was heated to 100 °C for 30 min in a digestion block. After cooling the hydrolysed solution was centrifuged at 2695 × g for 3 min and the clear supernatant decanted into a fresh glass tube. The residue was washed twice with 15 ml of deionised water and the washings added to the hydrolysate and kept in glass bottles at 4 °C until analysis for C and N using a TOC-V-TN analyzer (Shimadzu Corp., Kyoto, Japan).

To extract a less labile part of the soil C (fraction 2), the unhydrolysed residues were transferred to Pyrex tubes and dried at 60 °C. After cooling, 2 ml of 13 M H₂SO₄ was added, and the tubes were shaken overnight on a horizontal shaker at a speed of 80 strokes min⁻¹ at room temperature. Thereafter, 26 ml of deionised water was added to dilute the acid to 1 M and the residues were hydrolysed for 3 h at 100 °C with occasional shaking. After centrifugation at 2695 × g for 3 minutes, the clear hydrolysate was removed. The residues were washed twice with distilled water, and
the washings added to the hydrolysate and stored at 4 °C until analysis of C and N as described above. The remaining residual C was fraction 3 and was calculated by deducting the summed C fractions 1 and 2 from total organic C content of the soil (Belay-Tedla et al., 2009).

Acid hydrolysis of plant material followed the same protocol used for soils except that the sample size was decreased to 25 mg (Shirato and Yokozawa, 2006). The plant biomass C: acid ratio was the same as that used in the soil hydrolysis. As the residues could not be removed by centrifugation, the extracts were separated from un-hydrolysed residues by filtration using Whatman No. 1 filter papers (GE Healthcare UK Ltd.). After each hydrolysis, residues were washed twice with distilled water. Both soil and plant samples were analysed in triplicate.

2.7 Estimation of C pools throughout the soil profile

The total C stock and the absolute quantity of labile and recalcitrant C in different soil layers were estimated on area basis using bulk density determination of the different soil layers after adjustment for stone content. The total C pool size in soil profile (0-100 cm) was calculated by fitting a 2nd order quadratic to the soil C data; predicted values were then used to interpolate the C content of all soil layers in 10 cm increments. The actual and in-filled values were then summed to determine total size of the C pool.

2.8 Leaf litter decomposition

We studied the decomposition dynamics of leaf litter in each plots using leaf litter from respective species to examine the impacts of species identity or mixture on decay rates that affects the quality and quantity of SOC. We used leaves with a natural
water content rather than air dried leaves, because firstly, most of the leaves reach the forest floor as fresh litter in this ecosystem and thus we mimicked the natural process and secondly, air drying can substantially depressed the initial decay rates, especially in case of *A. glutinosa* (Taylor, 1998).

Litter decomposition rates were determined by mass loss of leaves in 180 nylon mesh bags. Fifteen litter bags (1 mm mesh, 20 cm × 15 cm), containing 5.0 g litter each, were placed on the forest floor of each tree species monoculture and the three tree species polyculture plot in July 2009. The bags were deployed at close contact with mineral soils under the litter layer to include the interaction of soil fauna activities, especially as the activity of earthworms at the site is high (Scullion et al., 2014). Litter representing the three species polyculture plots was composed of *B. pendula*, *A. glutinosa* and *F. sylvatica* in the ratios of 4:5:1 based on species contributions to litter fall baskets within the polyculture plots (Ahmed, 2011). Three litter bags were harvested after 3, 6, 10, 15 and 21 weeks from each plot. The litter was cleaned to remove soil particles, and dried at 60 °C for 72 h before weighing. A sub sample of 0.5-1.0 g was burned at 450 °C overnight, and weighed to determine ash content. Ash weight was deducted from the total litter weight to account for the contribution of adhered mineral soil particles to litter mass. The following single exponential decay model (Equation 1; Olson, 1963) was used fitted to the leaf mass loss data to compare leaf litter decay between species, where *A* is the initial litter mass, *k* is the decay rate constant, and *t* is time.

\[
\text{mass remaining} = Ae^{-kt}
\]

To determine the effect of litter mixture on litter decomposition, the mass loss of the polyculture litter bags was compared with those of a theoretical polyculture calculated from the mass loss of the single species litter bags. Equation 2 shows the
theoretical mixture biomass calculation, where $M_{\text{species}}$ is the mass contributing towards the mixture. The theoretical basis of this calculation is directly analogous to the Relative Yield of Mixtures index used to quantify the effects of competition (Wilson, 1988). The use of Equation 2 in this experiment is comparable with the Relative Yield Total (Weigelt and Jolliffe, 2003).

$$M_{\text{mixture}} = \left( \frac{4}{10} \times M_{\text{Betula}} \right) + \left( \frac{5}{10} \times M_{\text{Alnus}} \right) + \left( \frac{1}{10} \times M_{\text{Fagus}} \right)$$

Equation 2

2.9 Statistical analysis

The BangorDIVERSE experiment was designed as a fully replicated ($n=4$) field experiment with seven tree species planted in monoculture and mixtures of two and three species. For this research, three replicate plots each of grass, a single-tree species or a three tree species polyculture were studied. C pools were compared across four depths and four species types separately using One-way ANOVA, and pairwise comparisons made with Tukey’s HSD post hoc test (SPSS v14.0, SPSS Inc., Chicago, IL, USA). Normality was assessed by Shapiro-Wilk test and homogeneity of variances was determined by Levene’s test. Main effects were considered to be significant at $P<0.05$.

3. Results

3.1 Soil carbon
Total soil C content decreased with increasing depth in soils under all species types and the grassland (Fig. 2). Significant differences in soil C between *F. sylvatica* plots compared to *B. pendula* plots was observed at top two soil layers, but no differences in soil C were found in any of the treatments in soil below 20 cm. Soil microbial biomass C ranged between 0.56 ± 0.03 mg C kg\(^{-1}\) and 0.83 ± 0.08 mg C kg\(^{-1}\) in the top 0-10 cm of soil, and the species composition had no significant impact on the microbial biomass C and N content. In all the plots, the C:N ratio decreased with increasing soil depth (Suppl. Fig. 1), but no statistically significant differences between any of the treatments at any soil depth were observed. However, the greatest change in C:N ratio between the top and bottom soil layers was found in the grassland and the *B. pendula* plots (Δ 6.1) compared to a change of ca. 3.5 in the *A. glutinosa*, *F. sylvatica* and polyculture plots.

### 3.2 Relative contribution of C fractions to total C concentration

The C content of each of the SOM fractions varied between species and between soil depths. Figure 3 illustrates the distribution pattern of three C fractions in soils under different plant species and polyculture. In the 0-10 cm layer, fraction 1 was between 22 – 36 % of the total C, this increased to 29 – 52 % at 100 cm depth. The C of fraction 1 in the *F. sylvatica* plots was significantly (*P*=0.012 & *P*=0.002) greater than the soils of *A. glutinosa* plots at top two layers, and at 10-20 cm depth *B. pendula* and the polyculture soils were significantly higher than *A. glutinosa* (*B. pendula*, *P*=0.006; polyculture, *P*=0.035). For fraction 2, except in the grassland plots, the changes in percentage contribution to the total soil C were less pronounced compared to fraction 1. Grassland contained a significantly higher percentage of C in fraction 2 in all soil layers than other plots, except *A. glutinosa*. 
Fraction 3, the residual C after extraction, representing potentially the most recalcitrant C was unaffected by soil depth in *F. sylvatica* and the polyculture, but was significantly lower in the middle layer (40-50 cm) compared to the upper layers in the *B. pendula* and *A. glutinosa* plots. Species identity and mixture did not significantly affect the relative contribution of fraction 3 in the top two layers of soil (Fig. 3). Further down the soil profile, a significantly higher percentage of fraction 3 was found in polyculture soils than in *B. pendula* (*P*=0.014) and *A. glutinosa* (*P*=0.002) soils at 40-50 cm; and in the 100 cm layer the contribution was higher compared to *A. glutinosa* (*P*=0.013). In both the 40-50 and 100 cm layers, *A. glutinosa* had the lowest proportion of C within fraction 3 (27 and 34%, respectively).

### 3.3 Total C storage and C pool size of each fraction

The total C storage to a depth of 100 cm in the various plots ranged between 10.2 ± 0.9 under grass to 6.9 ± 0.8 kg C m⁻² under *F. sylvatica*, with no significant variation between the treatments (Table 1). In Table 1 the pools of C are shown as the total extractable (fraction 1 and fraction 2) and the residual C in fraction 3. The tree species grown in monoculture and polyculture showed no significant difference in total C stocks. We examined the influence of tree species on fraction 3 in upper (0-40 cm) and lower (40-100 cm) region of the soil profile (Fig. 4). In the upper layers no significant differences were found between the treatments, however in the deeper soil layers, the greatest storage of C in fraction 3 was found in the polyculture. The C storage in the polyculture soil at depth was significantly greater compared to the *B. pendula*, *A. glutinosa* and grassland soil, but not statistically different compared to *F. sylvatica*. Both *F. sylvatica* and the polyculture in the lower soil profile had a
significantly higher ($P=0.015$ and $P<0.001$) C storage in fraction 3 compared to the profile under grass.

3.4 Fractionation of litter C inputs

Total C in the leaves and roots of the three tree species was 52 and 53 %, respectively, significantly greater than the sward comprising the grassland which contained 44 % ($P=0.020$) and 40 % ($P=0.002$) for leaves and roots, respectively. In the tree leaves, C extracted from fractions 1 and 2, was similar. In contrast, in the roots of *F. sylvatica* the C content of fraction 2 was higher than in the other tree species (Fig. 5). In grass leaves and roots, the highest amount of C was in fraction 1, and the amount of fraction 3 was only 35 and 37 % of the total C, for leaves and roots, respectively.

3.5 Leaf litter decomposition

During the course of decomposition, mass remaining in leaf litter best fitted a first order exponential decay model. Decay rate coefficients for the three species grown in monoculture and polyculture are shown in (Table 2). Overall, and during the first four sampling intervals (3, 6, 10 & 15 weeks), there was a significant difference in mass loss between the litter of tree species ($P<0.001$; Table 2). During this period the rate of mass loss of the single species trees litter was highest in *A. glutinosa*, which was 1.94 and 1.80 times faster than *F. sylvatica* and *B. pendula*, respectively (Table 2). In the mixed species litter bags there was a dramatic and significant ($P<0.001$) reduction in mass loss, which was 4.36 times slower than *A. glutinosa* in monoculture.

4. Discussion
4.1 Tree traits

The storage and the distribution of organic C in soils are influenced by the quality and quantity of inputs determined by the integrated effects of species-specific traits (Schmidt et al., 2011). In this study, we examined three tree species selected due to their strongly contrasting productivity and functional traits, to accentuate the species-specific contribution to soil C pools. As a consequence of the trait differences, the species have different qualities of leaf litter inputs (see below), but also different rates of fine root turnover and hence root litter inputs (Smith et al., 2013b). The differences in leaf litter quality were reflected in the initial rates of decomposition, where the decomposition of *A. glutinosa* was nearly two times faster than the other two species. We found that decomposition processes of mixed species litter bags were slower than single species when deployed at our field site. Species-specific interactions during litter decomposition have been shown to have no effect (Prescott et al., 2000), retard (Chapman et al., 1988), or enhance decomposition processes (de Marco et al., 2011). We attribute the reduction in decomposition rates to the combination of a highly recalcitrant lignocellulose matrix of *F. sylvatica* litter and species-specific secondary metabolites, such as polyphenols and monoterpenes that inhibit N mineralisation and species-specific decomposer communities (Hattenschwiler et al., 2005). In addition to the potential interaction of late successional species litter chemistry in decomposition, and consistent with the findings of Giertych et al. (2006), we found a higher water-soluble polyphenolic content in *B. pendula* (20.5 mg L$^{-1}$) compared to *A. glutinosa* (17.5 mg L$^{-1}$) litter (Ahmed, 2011). The nitrogen content of senesced leaf litter was 30.5, 29.0 and 35.0 g kg$^{-1}$ for *F. sylvatica*, *B. pendula* and *A. glutinosa*, respectively, suggesting that the rapid initial decomposition of *A. glutinosa* was driven by nitrogen
availability. However, lignin content did not follow the same species order and was 138, 272 and 338 g kg\(^{-1}\) for \textit{A. glutinosa}, \textit{B. pendula} and \textit{F. sylvatica}, respectively, potentially leading to slower decomposition of fraction 3 C for \textit{F. sylvatica} relative to the other species.

Plant species identity can also influence the production and distribution of fine root biomass throughout the soil profile. The effect of species diversity on root biomass and production is extremely variable with studies showing no effect (Bauhus et al., 2000), a reduction (Bolte and Villanueva, 2006) or increase (Brassard et al., 2011). In the species used here, \textit{A. glutinosa} had the highest rate of fine root turnover, and \textit{F. sylvatica} the highest fine root length in the top 30 cm of soil (Smith et al., 2013b). However, differences in the rate of fine root turnover were not seen in all years (Ahmed, 2011). In addition to the influence of plant litter chemistry and species identity on decomposition, the phenology of leaf and root growth can also influence belowground processes (Niinemets and Tamm, 2005). Indeed, seasonality has a particularly strong control on the phenology of grassland species (Steinaker and Willson, 2008).

### 4.2 Organic C storage of soils under different plant species

No significance difference in SOC stock (0-100 cm) was observed both between the tree species, and in comparison to the grassland (Table 1). This is in consistent with the study of Vesterdal et al. (2008), who reported no significant variation in the soil C stocks of five European broadleaved tree species, including \textit{F. sylvatica}, after 30 years of growth. However, we did find a higher organic C concentration in \textit{B. pendula} soil compared to \textit{F. sylvatica} in the upper two layers of the soil profile (Fig. 2). The biomass production and subsequent litter fall in \textit{B. pendula} was much higher than the
late successional species *F. sylvatica*, and may be the cause of the higher SOC in the upper soil layers.

4.3 Soil organic carbon and fractionation

Changes in total soil C were only observed in the top 20 cm of soil, below this depth soil C was not different between the tree species or the grassland. Soil C stock change and physical fractionation were investigated at the Bangor site using different but adjacent plots with the same species by Hoosbeek et al., (2011), who found an increase of 530 g C m$^{-2}$ in the top 0-10 cm layer 4 years after planting. In contrast to our chemical fractionation results, the distribution of coarse, fine and aggregate particulates were similar between all species and the polyculture. Here we showed that using chemical fractionation, the organic C content of fraction 1 was significantly lower in soils under *A. glutinosa* than in *F. sylvatica* in the 0-10 cm soil layer (Fig. 3).

We propose two mechanisms to explain our observed differences in fraction 1 C content. First, relative to *F. sylvatica*, *A. glutinosa* is poor at translocating nutrients and carbohydrates during senescence (Lecerf and Chauvet, 2008), and subsequently the senesced litter of *A. glutinosa* is considered to be of high quality due to thick, mesophyll rich, leaves with a low C:N ratio. These traits, which are favourable to grazing by soil organisms and microbial decomposition (Kazakou et al., 2009), probably resulted in a rapid removal of fraction 1. The influence of litter quality on decomposition processes was also supported by significantly faster litter mass loss of *A. glutinosa* than *F. sylvatica* during the first three weeks (Table 2). Second, during organic matter mineralization and microbial turnover, C not respired as CO$_2$ is retained within microbial biomass, or released as dissolved organic carbon (DOC),
which then leaches through the soil profile reducing the size of the pool in shallow soil layers (Currie and Aber, 1997).

The percentage of C in fraction 1 increased gradually down the soil profile under *B. pendula* and *A. glutinosa*, but remained constant under *F. sylvatica* and the polyculture. Studies using stable C isotopes and radiocarbon have revealed that acid hydrolysable C, as in fraction 1, as well as mineral associated C are consistently younger than other fractions (Leavitt et al., 1997), and generally, it is assumed that the age of soil C increases with depth (Fontaine et al., 2007). Potential sources of the fraction 1 in the lower soil layers are numerous and include translocation from upper soil layers with DOC or bioturbation, root exudates (de Graaff et al., 2010), and priming of more recalcitrant soil organic matter (Rovira and Vallejo, 2007, Kogel-Knabner et al. 1991) with subsequent release of more labile fractions. Earthworm activity at the site is high, and the earthworms were shown to have a higher preference for litter from *B. pendula* and *A. glutinosa* than that from *F. sylvatica* (Scullion et al. 2014).

The largest percentage of C in the top soil layer was in fraction 3, and was not influenced by tree species identity or grass. This is consistent with the findings of Hoosbeek et al., (2011) who reported that the physical fractionation of particulate organic matter at the same experiment site and found that tree species identity had no effect on soil C stabilization processes and microaggregate protection in the upper soil layers. A caveat of our acid hydrolysis approach to separating C fractions is that physically protected labile C may be included in fraction 1 (McLauchlan and Hobbie, 2004), but there seems to be broad agreement between the acid hydrolysis and physical fractionation methods. Similarly, it has also recently be shown that using acid fractionation schemes more aggressive than the one used here, can lead to de-
novo synthesis of non-hydrolysable substances and loss of pure model carbohydrates (Greenfield et al. 2013).

Soils from under grass differed strongly to the tree plots in the distribution of all three C fractions. Throughout the soil profile, the strong difference between trees and grass were most likely attributable to a shallow rooting depth and turnover of non-woody grass roots. In addition, the higher amount of fraction 1 C in grassland soil might be due to the high quality of grass litter, which contains substantially less lignin than tree litter, and thus is easily decomposed (Deschaseaux and Ponge, 2001). This is supported by our analysis of grass leaf and root material, which showed much higher quantities of fraction 1 C in grass than in tree materials (Fig. 5).

4.4 Recalcitrant C in deep soil layers

There were no differences in the percentage of soil C found at 40-100 cm soil depth between the treatments. Below 40 cm the percentage soil C was less than 1 %. However, surprisingly, significantly greater quantities of the total soil C were found in fraction 3 in the three species tree polyculture stands, compared to the single species stands of B. pendula and A. glutinosa and the grassland. The higher fraction 3 storage was found in all three replicate polyculture plots irrespective of previous land-use (Fig. 1), as was the lower fraction 3 storage in the grassland plots. Similarly, in the polyculture plots, there were no obvious differences in soil texture, such as high levels of clay, which could account for the increased fraction 3 storage. The higher fraction 3 storage in polyculture is difficult to reconcile with either C inputs via above- or below-ground biomass of the different species. By far the biggest difference in fraction 3 storage, or the percentage of fraction 3 in the total soil C, was found when the trees were compared to the grassland. This suggests that the differences in fraction
3 storage could be related to depth distribution and timing of C inputs, litter quality, or quantity. A potential mechanism could be a priming effect, where the input of labile C products into the deep soil layers, possibly via the flow of DOC stimulated microbial mineralisation of old C (Kuzyakov et al., 2000; Hoosbeek et al., 2004). Microbial priming of deep soil C was reported by Fontaine et al., (2004) who demonstrated that fresh inputs of labile C allowed the co-metabolism of old recalcitrant C by the microbial community at a depth of 60-80 cm. Therefore, the availability of easily biodegradable compounds through vertical DOC transport into deep soil layers could be an important factor in mediating the storage of recalcitrant C in deep soil layers. The chemical composition of DOC is related to the plant litter from which it is derived (Hansson et al., 2010). Our leaf litter decomposition experiment showed a large and significant difference between the rate of decomposition in monoculture and the three species polyculture, which may have resulted in a greater amount of recently derived C moving down the soil profile and microbial priming fraction 3 at depth. Again difficult to reconcile with current ideas is the speed at which these changes must have occurred. However, it must be stressed that the levels of total C at 100 cm soil depth are very low, enabling detection of small changes.

To conclude, our data suggest that even within relatively short time scales vegetation types and tree species identity and mixtures can influence both accumulation of soil C in surface layers, but also the storage of more recalcitrant fractions in deeper soil layers. This may be due to the direct inputs of new C, but also due to the effects of new C influencing the levels and distribution of old soil C.

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References


Ahmed, I.U. 2006. Leaf decomposition of birch (Betula pendula), alder (Alnus glutinosa) and beech (Fagus sylvatica) grown under elevated atmospheric CO2. Master thesis, SENRGY, Bangor University, Bangor, UK.


microbial carbon and nitrogen pools of a tall grass prairie soil in the US Great
Plains subjected to experimental warming and clipping. Soil Biol. Biochem.,
41, 110–116.

Binkley, D., Valentine, D. 1991. Fifty-year biogeochemical effects of green ash, white
pine, and Norway spruce in a replicated experiment. For. Ecol. Manage., 40,
13–25.

Bolte, A., Villanueva, I. 2006. Interspecific competition impacts on themorphology
and distribution of fine roots in European beech (Fagus sylvatica L.) and

productivity between mixed- and single-species stands. Func. Ecol., 25, 238–
246.

Produktivität des Wasserverbrauches bei Holzpflanzen. Allg Forst-und Jagd-
Ztg 145, 81–86.

Carnus, J., Parrotta, J.A., Brockerhoff, E., Arbez, M., Jactel, H., Kremer, A., Lamb,
D., O’Hara, K., Walters, B. 2006. Planted forests and biodiversity. J. For., 104,
65–77.

Carvalheiro, K.D., Nepstad, D.C. 1996. Deep soil heterogeneity and fine root
distribution in forests and pastures of eastern Amazonia. Plant Soil, 182, 279–
285.

litters of tree mixtures compared with pure stands. Agric. Ecosys. Environ.,
24, 33–40.


Fontaine, S., Bardoux, G., Abbadie, L., Mariotti, A. 2004. Carbon input to soil may

Giertych, M.J., Karolewski, P., Zytkowski, R., Oleksyn, J. 2006. Differences in
defence strategies against herbivores between two pioneer tree species, *Alnus*

Greenfield, L. G., Gregorich, E. G., van Kessel, C., Baldock, J. A., Beare, M. H.,
Billings, S. A., Clinton, P. W., Condron, L. M., Hill, S., Hopkins, D. W.,
Janzen, H. H. 2013. Acid hydrolysis to define a biologically-resistant pool is
compromised by carbon loss and transformation. Soil Biol. Biochem., 64,
122-126.

Hagen-Thorn, A., Callesen, I., Armolaitis, K., Nihlgard, B. 2004. The impact of six
European tree species on the chemistry of mineral topsoil in forest plantations

Hansen, K., Vesterdal, L., Schmidt, I.K., Gundersen, P., Sevel, L., Bastrup-Birk, A.,
Pedersen, L.B., Bille-Hansen, J. 2009. Litterfall and nutrient return in five tree
species in a common garden experiment. For. Ecol. Manage., 257, 2133–2144

mineralised and leached as DOC during decomposition of Norway spruce

Hattenschwiler, S. 2005. Effects of tree species diversity on litter quality and
decomposition. In: Scherer-Lorenzen M, Korner CH, Schulze ED (eds) Forest
diversity and functions, temperate and boreal systems. Ecological Studies 176.
Springer-Verlag, Berlin, pp 149.


Figure Legends

**Fig. 1** Plot positions overlayed on to a krigged plot of initial soil organic matter content determined in the top 0-10 cm layer on a 10 × 10 m grid across the two fields of the BangorDIVERSE experimental site. In field 1 the gray overlay marks the approximate extend of the short rotation coppice trial. Coloured boxes represent plots of *A. glutinosa* (green), *B. pendula* (yellow) and *F. sylvatica* (blue) in monoculture, and a three species tree polyculture (orange), or a grassland (red).

**Fig. 2** Vertical distribution of soil organic carbon from under monoculture or polyculture stands of *B. pendula*, *A. glutinosa*, and *F. sylvatica*, or a grassland. Symbols show means ± SE (*n*=3), statistically significant differences (*P*<0.05) are denoted by a superscript asterisk.

**Fig. 3** The contribution of fractions 1, 2 and the residual fraction 3 to the total C pool in different soil layers from under monoculture and a three species polyculture of *B. pendula*, *A. glutinosa* and *F. sylvatica*, or a grassland. Shown are means ± SE (*n*=3), statistically significant differences are denoted by a superscript asterisk (*P*<0.05, **P**<0.01 and ***P**<0.001).

**Fig. 4** The total soil pool (kg C m$^{-2}$) of fraction 3 as determined by sequential acid extraction in a 1 m deep soil profile under *B. pendula*, *A. glutinosa*, *F.*
*sylvatica* and grassland. Bars show means ±SE (n=3). Bars not followed by similar indices are statistically significant (P<0.05).

**Fig. 5** C fractions in leaves and roots of *B. pendula*, *A. glutinosa*, *F. sylvatica* and a grassland as determined by sequential acid hydrolysis. Values shown are expressed as the percentage of total C. Shown are means ± SE (n=4). Bars with same indices are not statistically significant (P<0.05).

**Suppl. Fig. 1** Relationships between clay content and (a) total soil organic C, (b) labile C fraction 1 and (c) recalcitrant C (fraction 3) for soils under monoculture and a three species polyculture of *B. pendula*, *A. glutinosa* and *F. sylvatica*, or a grassland.