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Article

Variety and Site Drive *Salix* Mixture Effects on Soil Organic Matter Chemistry and Soil Carbon Accumulation

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Abstract: Soil organic matter (SOM) is essential for nutrient cycling and soil carbon (C) accumulation, both of which are heavily influenced by the quality and quantity of plant litter. Since SOM dynamics in relation to plant diversity are poorly understood we investigated the effects of willow variety and mixture, and site on the soil C stocks, SOM chemical composition and thermal stability. Using pyrolysis-field ionization mass spectrometry we analysed the top 10 cm of soil from two 7-year-old experimental sites in Germany and Sweden, with monocultures and mixtures of two willow varieties (*Salix* spp.) belonging to different species. Overall, site had the strongest effect on SOM quality. Results showed significant variability across sites for willow identity and mixture effects on C accumulation and SOM chemistry. In the German site (Rostock), yearly soil C accumulation was higher ($p < 0.05$) for variety 'Loden' ($1.0 \text{ Mg C ha}^{-1} \text{ year}^{-1}$) compared to 'Tora' ($0.5 \text{ Mg C ha}^{-1} \text{ year}^{-1}$), whilst in the Swedish site (Uppsala), both varieties exhibited similar soil C accumulation rates of around $0.6 \text{ Mg C ha}^{-1} \text{ year}^{-1}$. Willow variety identity significantly affected SOM quality at both sites, while mixing had minor effects. Our findings emphasize the significance of site-specific context and variety or species identity in shaping soil C accumulation in willow plantations.

Keywords: variety or species mixing; Soil organic matter; carbon sequestration; chemical composition; thermal stability

1. Introduction

Soils store more carbon (C) than both the atmosphere and all vegetation combined [1], thereby playing a critical role in terrestrial ecosystem CO₂ exchange [2], and in balancing atmospheric CO₂ emissions. Whether soils act as sinks or sources of CO₂ depends on the balance between photosynthesis, respiration and stabilization of C belowground, and changes in soil C stocks have the potential to impact atmospheric CO₂ levels and the global C budget [3]. Soil organic matter (SOM) comprises the largest active reservoir of terrestrial organic carbon [4], and plays a direct role in climate change mitigation [5–7]. However, the dynamics of SOM are complex and not fully understood [8], and improved understanding of its formation and stabilization is crucial for developing sustainable approaches to enhance soil C sequestration.

The SOM consists of decomposing plant material and microbial-derived compounds varying in size, extent of degradation, accessibility, and residence time [9]. The chemical composition and stability of SOM controls the rate at which microbial communities process organic matter, thus

governing the accumulation of soil C [10,11]. Since soil respiration is influenced by the soil microbial activity a higher soil respiration rate could therefore reflect increased decomposition rates. In the present study we use chemistry indicating SOM stability as a proxy of SOM quality. The stability of SOM is partly determined by its constituent molecules, some of which are more resistant to decomposition (e.g., lignin, phenols and suberin; [12,13]) than others (e.g., free fatty acids and peptides; [14,15]). Higher proportions of recalcitrant molecules contribute to more stable SOM, which is more likely to accumulate over time [16]. Climatic and edaphic factors further complicate SOM dynamics, as temperature and precipitation influence microbial activity and soil type, which can be instrumental for SOM stabilization [17–19].

There are many wet-chemical, spectrometric and spectroscopic analytical methods available to assess the quality of SOM, e.g., [20,21]. Pyrolysis-field ionization mass spectrometry (Py-FIMS) is a powerful analytical tool, with which biogenic marker substances can be associated with molecular compound classes derived from soil samples, thus providing a means of characterizing SOM chemical composition while also providing the possibility to assess the thermal stability of these same biogenic markers [11,22]. It reveals the strength of chemical bonds within molecules or between reactive mineral surfaces and organic molecules in response to thermal stress, thus providing a measure of resistance to microbial decomposition [11,13]. This data-rich method provides a high-resolution overview of the SOM chemical structure with the potential to elucidate responses of belowground processes to factors such as plant species identity and plant species mixing. SOM chemical composition and thermal stability are important aspects to consider when examining how aboveground vegetation influences the stability of SOM in different environments.

The quantity and quality of plant litter input can vary greatly between species and vegetation types [23,24]. Multiple studies have shown plant diversity to enhance soil C sequestration [25,26], often through increased aboveground biomass production through altered resource competition or facilitative plant-plant interactions, thus increasing the amount of organic matter supplied to the soil and therefore the accumulation of C [27–29]. However, effects of species diversity on plant productivity are often highly context dependent [30,31]. Further, an increase in C storage with plant diversity can also result from longer persistence of plant litter due to slower decomposition [32]. Plant diversity, mainly studied in grasslands, but with evidence emerging from forest ecosystems, has been shown to affect SOM chemical diversity [33–35], microbial composition, activity and biomass [36,37], soil nutrient retention [38], and soil gas emissions [39]. Increased soil microbial biomass and consequently later necromass can make up more than half of SOC [40]. Plant diversity should therefore likely influence both SOM chemical composition and thermal stability, which could have major implications for SOM decomposition and ultimately ecosystem functioning [41–44]. Yet, the effects of plant diversity are not thoroughly explored in forest ecosystems, and especially the responses in SOM chemical composition and stability to tree species mixing remain poorly understood.

The cultivation of *Salix* spp. (willows) as short rotation coppice (SRC) has emerged as a promising approach to sustainably produce renewable biomass [45,46], with the largest cultivated areas found in China and Argentina, followed by Europe [47]. These production systems are characterized by short growth cycles of 2–5 years, after which the stems are harvested and shoots regrow rapidly from the stumps left in the soil [47]. In addition, willow SRC can serve as a model system for investigating plant diversity-productivity relationships [48], due to the fast growth and high phenotypic variability of willow species along with moderate to small phylogenetic contrasts. This enables the evaluation of subtle phenotypic differences on plant-plant interactions [49,50]. Different willow species and varieties can have differential impact on belowground C dynamics and soil microbiota [51–53], and willow species or varieties grown in mixtures can differ from their monoculture counterparts in terms of nitrogen economy and productivity [50,54]. In the present study, we used two phenotypically distinct willow varieties belonging to different species grown both as monoculture stands and as mixtures. The aim of the study was to evaluate the effects of willow variety identity, mixture and site on soil C accumulation, SOM chemical composition and thermal stability using Py-FIMS methodology. We hypothesized that: (i) willow variety identity

affects soil C accumulation, SOM chemical composition and stability, (ii) willow variety mixtures accumulate similar amounts of soil C but differ in SOM chemical composition and stability compared to the equivalent monocultures at the same site, (iii) site conditions modulate the effects of variety identity and mixture on soil C accumulation, SOM chemical composition and stability.

2. Materials and Methods

2.1. Experimental Design and Plant Material

Two experimental sites, one in Rostock, Germany (54°02' N 12°05' E) and one in Uppsala, Sweden (59°49' N 17°39' E) were planted with willow in a short rotation coppice (SRC) system on former arable land in 2014. The two sites are part of the ECOLINK-Salix research trials as well as members of the global tree diversity network (TreeDivNet; [55]). Climatic conditions and soil properties differ between the two sites, notably clay content is roughly 10 times higher in the Uppsala site compared to the Rostock site (Table 1).

Table 1. Site characteristics: soil group [56], topsoil (0-10 cm depth) properties measured in 2021 and climatic conditions during time since establishment at sites Rostock and Uppsala.

Site	Soil group	pH	Bulk density [g cm ⁻³]	Clay content [%]	MAT [° C]	MAP [mg g ⁻¹]
Uppsala	Vertic Cambisol	5.2	1.4	52	7.53	500
Rostock	Stagnic Cambisol	6.2	1.3	5	10.35	730

Two phenotypically distinct willow varieties belonging to different *Salix* species were used as stand components: ‘Loden’ (L; *S. dasyclados* Wimm.) and ‘Tora’ (T; *S. schwerinii* x *S. viminalis*). Generally, higher shoot biomass production is reported for ‘Tora’ than for ‘Loden’ [50,54], while leaf and fine root biomass production has been found to be higher for ‘Loden’ than for ‘Tora’ [57,58]. ‘Loden’ and ‘Tora’ also differ in terms of leaf chemistry and element stoichiometry [50,59], with [60] reporting weak effects of site, strong effects of variety and interaction effects between site and variety, on leaf litter decomposability as measured by fraction of remaining biomass after incubation. Willows can form both ecto- and arbuscular mycorrhizal associations [61], but are generally more associated with ectomycorrhizal fungi [62]. In addition, [63] reported on greater ectomycorrhizal colonization of root tips belonging to *S. dasyclados* Wimm. than *S. viminalis* L., which are similar to ‘Loden’ (a *S. dasyclados* variety) and ‘Tora’ (a *S. schwerinii* x *S. viminalis* variety), respectively. In the present study we use ‘Loden’ and ‘Tora’ in monoculture and the 2-component mixture. The willow varieties were planted in a randomized block design with 3 replicates for each monoculture (‘Loden’ and ‘Tora’) and mixture (‘Loden’:‘Tora’) for each site (i.e., 9 plots per site). All 9.6 × 9.6 m plots were planted in a hexagonal pattern with 12 rows, each row containing 12 plants at a distance of 0.8 m between neighboring plants, and further details can be found in [64].

2.2. Soil Sampling, Analyses and Measurements

Since the majority of *Salix* fine roots are found within the top 10 cm of soil [65,66], we focused our sampling efforts on this layer, assuming the most significant impact of *Salix* growth occurs here due to the turnover of fine roots and leaf litter. We collected nine soil samples per plot per site at experiment establishment in April 2014 and during the third cutting cycle in April 2021, using a soil auger (3 cm diameter). Subsamples were pooled per plot, dried at 40 degrees for 48 h and sieved (< 2 mm). The concentrations of C and N were determined using a CN analyzer (Vario EL, Elementar Analysensysteme, Hanau, Germany). Soil pH was determined in 0.01 M CaCl₂ at a soil:solution ratio of 1:2.5 (w/v). The C stocks (Mg ha⁻¹) were calculated for 2014 and 2021 by the following formula:

$$C_{stocks} = C_{org} * \rho d * depth$$

(1)

where C_{org} is the organic C concentration ($\text{g } 100 \text{ g}^{-1}$) in the soil, ρd is the dry bulk density (g cm^{-3}), and $depth$ is the soil sampling depth (10 cm). Yearly soil C accumulation ($\text{Mg C ha}^{-1} \text{ year}^{-1}$) was calculated as the difference between 2021 and 2014 C stocks averaged per year. The determination of the dry bulk density (ρd) was based on the use of 250 cm^3 cores in 2014 and 2021 ($\rho d = \text{dry weight [g]} / \text{volume [} 250 \text{ cm}^3 \text{]}$). The soil cores were dried at $105 \text{ }^\circ\text{C}$ until a constant weight was achieved.

Double lactate-extractable phosphorus (P_{dl}), potassium (K_{dl}) and magnesium (Mg_{dl}) were determined after extraction of 0.6 g soil with 30 ml lactate solution [67]. The element concentrations in the extract were measured with inductively coupled plasma-optical emission spectroscopy (ICP-OES, Optima 8300, Perkin Elmer, Waltham, Massachusetts, USA). The lactate-extractable concentrations have been considered to represent the plant-available fraction in soils [68].

CO_2 respiration was measured at 4-week intervals (8 times) during the growing season (April to October) for year 2021 at the Uppsala site as a proxy for decomposition. Measurements were made using a portable infrared gas analyzer coupled to a 1296 cm^3 dark chamber in closed air circuit (EgM-4 with SRC-1 probe type; PP Systems, Amesbury, Massachusetts, USA). The chamber was pressed firmly to the ground (avoiding weeds) for measurements in at least four different places per plot at each sampling occasion, evenly spread out with the intention to reflect the plot-level treatments. Due to cracking clay soil surfaces during dry summer months the sampling place varied between sampling occasions, ensuring a tight seal and avoiding cracks in the soil where large amounts of CO_2 was released. Respiration was typically monitored between 9:00 and 14:00 CET on cloudy to half-cloudy days after a recent rainfall if possible for ca 120 s per measurement and calculated as a function of the linear increase of CO_2 concentration in the chamber. Measurement quality was ensured by automatic baseline calibration by the ‘auto-zero’ option at least every 20 min and by keeping the start CO_2 concentration in the chamber at ambient level [69].

2.4. Pyrolysis-Field Ionization Mass Spectrometry (Py-FIMS)

SOM chemical composition and thermal stability were analysed by pyrolysis-field ionization mass spectrometry (Py-FIMS) for samples from 2021 in three plots each for ‘Loden’, ‘Tora’ and ‘Loden’:‘Tora’ plots from the Rostock site and from two plots each from the Uppsala site. About 3 mg of the air-dried, ground and homogenized samples were thermally degraded by pyrolysis in the ion source (emitter: 4.7 kV, counter electrode -5.5 kV) of a double-focusing Finnigan MAT 95. The samples were heated in a vacuum of 10^{-4} Pa from $50 \text{ }^\circ\text{C}$ to $700 \text{ }^\circ\text{C}$, in temperature steps of $10 \text{ }^\circ\text{C}$ over a time period of 18 min. Between magnetic scans the emitter was flash heated to avoid residues of pyrolysis products. About 65 spectra were recorded for the mass range m/z 15 to 900. Biogenic marker signals (m/z) were assigned to nine relevant compound classes according to [21] (Table 2). All recorded marker signals (m/z) were combined to obtain the total ion intensity (TII) for each measurement. The difference in sample weight before and after pyrolysis provides a measure of “volatile matter” (VM) and is used to normalize sample ion intensities per mg sample weight. The hexoses:pentoses ratio is a measure of microbial- to plant-derived sugars [70] and is used as a measure of microbial contribution to SOM. Additionally, the ion intensities at each temperature step during pyrolysis was calculated separately for each of the about 65 single scans. The ion intensities of each compound class, plotted against the volatilization temperature, provided distinct thermograms that could be evaluated in terms of the thermal stability of compound classes.

Table 2. Pyrolysis-field ionization mass spectrometry (Py-FIMS) parameters and compound classes with associated biomolecules, as assigned by; [21,71–74].

Py-FIMS parameters	Explanations
Hexoses:pentoses	Ratio of microbial- to plant-derived sugars
TII	Total ion intensity ($10^6 \text{ counts mg}^{-1}$)
VM	Volatile matter in % ($\text{weight}_{\text{before pyrolysis}} / \text{weight}_{\text{after pyrolysis}}$)
CHYDR	Carbohydrates with pentose and hexose subunits
PHLM	Phenols and lignin monomers
LDIM	Lignin dimers

LIPID	Lipids, alkanes, alkenes, bound fatty acids, and alkylmonoesters
ALKYL	Alkylaromatics
NCOMP	Mainly heterocyclic N-containing compounds
PEPTI	Peptides (amino acids, peptides and aminosugars)
SUBER	Suberin
FATTY	Free fatty acids C ₁₆ -C ₃₄

2.5. Statistical Analyses

All statistical analyses were conducted in R (version 4.3.0; [75]). SOM chemical composition for the two willow varieties in monoculture and mixture per study site was visualized using principal component analysis (PCA; function *rda* in package *vegan*; [76]; Figure 2). The PCA was performed on the relative ion intensities for the 9 Py-FIMS compound classes (Table 2). We used a principal component approach for the compound class data due to significant covariance between several of the compound classes. Principal component scores (PC1 and PC2; function *scores* in package *vegan*; [76]) were used as response variables in mixed effects principal component regression models [77] to assess multivariate differences in SOM chemical composition between treatments.

The Shannon diversity index (*H'*), which accounts for both presence and relative abundance, was applied to our Py-FIMS biogenic marker signals (*m/z*) to quantify the chemical diversity of SOM (function *diversity* in package *vegan*; [76]):

$$H' = - \sum_{i=1}^n p_i \ln(p_i) \tag{2}$$

where *p_i* is the proportion of relative ion intensity for marker signal (*m/z*) *i*, and *n* is the number of measured marker signals in a given sample.

The SOM thermal stability, here used as an indicator of resistance to microbial decomposition [21], was calculated for total ion intensity (TII) as well as for each compound class separately (Table 2). This was done following [78] by dividing the sum of ion intensities volatilized at high temperature (> 400 °C) by ion intensities volatilized over the whole temperature range (50-650 °C).

Net diversity effects (NDE) of selected SOM properties were evaluated following [27]:

$$NDE \text{ (net diversity effect)} = \frac{Observed-Expected}{Expected} \tag{3}$$

in which, the observed values under variety mixtures are compared to the expected mixture values, calculated as the average values under their monoculture counterparts. Mixture effects are characterized as non-additive if significantly different from zero (NDE ≠ 0) or additive if they are not (NDE = 0). To avoid any mixture effects being occluded by potential dominance effects, we applied weights to the expected mixture values based on their proportional basal area in the mixtures [29] (Supplementary Material Table S1).

Responses in soil C accumulation, nutrient concentrations, Py-FIMS compound classes and SOM thermal stability were modeled using mixed effects multiple linear regression models with the fixed effects of variety composition ('Loden', 'Loden':'Tora' and 'Tora'), site (Rostock and Uppsala) and their interaction (function *lme* in package *nlme*; [79]). Block nested in site was added to the model as a random effect to account for site-specific block effects, which were especially noticeable at the Rostock site. Due to different variance structures at our sites, site-specific variance weights were included in the model. Soil CO₂ respiration was sampled only in the Uppsala site and was modeled using a mixed effects multiple linear regression model in response to the fixed effects of variety composition and undergrowth ground cover (%). Block and plot nested in block were added as random effects to account for block effects and repeated measurements respectively. Temporal autocorrelation between months was accounted for using an autoregressive correlation structure. For each response variable in the papaer, the statistical significance of site, variety composition and their interaction was evaluated using analyses of variance (ANOVA; type = III, Kenward-Roger's method, function *anova* in base R) and the results of these analyses are presented in Supplementary Material

Tables S2–S7. This was followed by pairwise comparisons of estimated marginal means using a Tukey adjustment (function *emmeans* in package *emmeans*; [80]). The function *contrast* from the *emmeans* package was used to examine specific variations across sites, varieties within sites, varieties across sites, as well as the disparities between expected and observed mixture values within sites and observed mixture values across sites. Estimated marginal means were used to calculate mean values due to the partially unbalanced nature of our dataset.

3. Results

3.1. Site Characteristics

The contents of soil C ($p = 0.004$), N ($p = 0.001$), K_{dl} ($p = 0.002$) and Mg_{dl} ($p = 0.011$) were significantly lower at the Rostock than Uppsala site (Table 3), while C:N ratio and P_{dl} were similar between sites. No differences in the above characteristics were found between willow varieties grown in monoculture or mixture at either site.

Table 3. Soil chemical characteristics and soil respiration: estimated marginal means of topsoil (0-10 cm depth) soil C:N ratios, nutrient concentrations of C, N, K_{dl} , Mg_{dl} and P_{dl} as well as soil CO₂ respiration for varieties ‘Loden’ and ‘Tora’ grown in monoculture and mixture at sites Rostock and Uppsala. Different letters indicate significant differences across sites (x-y) and within each site (a-b; $p < 0.05$, Tukey’s post hoc test).

Site	Variety composition	C:N	C [%]	N [%]	K_{dl} [mg g ⁻¹]	Mg_{dl} [mg g ⁻¹]	P_{dl} [mg g ⁻¹]	CO ₂ [g C m ⁻² h ⁻¹]
Rostock	‘Loden’	10.80 ^{xa}	1.26 ^{ya}	0.12 ^{ya}	9.99 ^{ya}	20.42 ^{ya}	4.40 ^{xa}	-
	‘Loden’:‘Tora’	10.63 ^{xa}	1.22 ^{ya}	0.11 ^{ya}	10.22 ^{ya}	21.06 ^{ya}	4.40 ^{xa}	-
	‘Tora’	10.22 ^{xa}	1.01 ^{ya}	0.10 ^{ya}	10.71 ^{ya}	21.96 ^{ya}	4.02 ^{xa}	-
Uppsala	‘Loden’	10.69 ^{xa}	1.89 ^{xa}	0.18 ^{xa}	22.23 ^{xa}	32.10 ^{xa}	4.27 ^{xa}	0.493 ^a
	‘Loden’:‘Tora’	10.58 ^{xa}	1.58 ^{xa}	0.16 ^{xa}	23.62 ^{xa}	27.94 ^{xa}	5.04 ^{xa}	0.453 ^a
	‘Tora’	10.85 ^{xa}	1.90 ^{xa}	0.17 ^{xa}	19.89 ^{xa}	28.76 ^{xa}	3.60 ^{xa}	0.461 ^a

The total thermal stability of bulk SOM was similar across sites, but total ion intensity (TII; $p = 0.004$, hexoses:pentoses ratio ($p = 0.004$) and chemical diversity (H' ; $p < 0.001$) were all significantly higher in Rostock than Uppsala (Table 4). Volatile matter was significantly higher in Uppsala than Rostock (VM; $p = 0.003$). In terms of compound classes, the Uppsala site had significantly higher relative abundances of carbohydrates (CHYDR; 5.0% TII, $p < 0.001$), phenols and lignin monomers (PHLM; 5.9% TII, $p < 0.001$), alkylaromatics (ALKYL; 3.5% TII, $p < 0.001$), N-compounds (NCOMP; 2.2% TII, $p < 0.001$) and peptides (PEPTI; 2.3% TII, $p = 0.001$); while lower relative abundances of lipids (LIPID; -1.7% TII, $p = 0.003$), suberin (SUBER; -0.2% TII, $p < 0.001$) and free fatty acids (FATTY; -0.5% TII, $p = 0.015$) compared to the Rostock site.

Table 4. Py-FIMS parameters: estimated marginal means of TII (total ion intensity; 106 counts mg⁻¹) and total thermal stability of bulk SOM (ions volatilized > 400 °C / ions volatilized 50-650 °C), VM (volatile matter), H' (SOM chemical diversity) and hexoses:pentoses (ratio of microbial- to plant-derived sugars) by site and by variety composition. Different letters indicate significant differences across sites (x-y) and within each site (a-b; $p < 0.05$, Tukey’s post hoc test).

Site	Variety composition	TII [10 ⁶ counts mg ⁻¹]	Total thermal stability	Chemical diversity [H']	Hexoses: pentoses	Volatile matter
Rostock	‘Loden’	53.3 ^{xa}	0.68 ^{xa}	6.71 ^{ya}	5.79 ^{xa}	2.83 ^{xa}
	‘Loden’:‘Tora’	39.1 ^{xa}	0.72 ^{xa}	6.30 ^{ya}	5.73 ^{xab}	2.92 ^{xa}
	‘Tora’	34.8 ^{xa}	0.77 ^{xa}	5.82 ^{ya}	5.67 ^{xb}	2.75 ^{xa}
Uppsala	‘Loden’	14.8 ^{ya}	0.84 ^{xa}	14.66 ^{xa}	5.47 ^{ya}	1.89 ^{yb}

	'Loden': 'Tora'	14.2 ^{ya}	0.82 ^{xb}	9.69 ^{xb}	5.45 ^{ya}	1.91 ^{yb}
	'Tora'	24.7 ^{ya}	0.83 ^{xab}	13.36 ^{xa}	5.53 ^{ya}	2.28 ^{ya}

3.2. Effects of Willow Variety

Soil C stocks measured at the establishment of the experiment in 2014 were similar across all plots within each site and averaged at 10.32 and 20.54 Mg C ha⁻¹ for sites Rostock and Uppsala, respectively. Seven years after planting, yearly soil C accumulation rates measured at the Rostock site under 'Loden' (1.02 Mg C ha⁻¹ year⁻¹) were significantly higher ($p = 0.004$) than under 'Tora' (0.54 Mg C ha⁻¹ year⁻¹). Soil C accumulation rates measured at the Uppsala site were similar between 'Loden' (0.55 Mg C ha⁻¹ year⁻¹) and 'Tora' (0.65 Mg C ha⁻¹ year⁻¹). Furthermore, neither Loden nor Tora differ in C accumulation rates between sites (Figure 1).

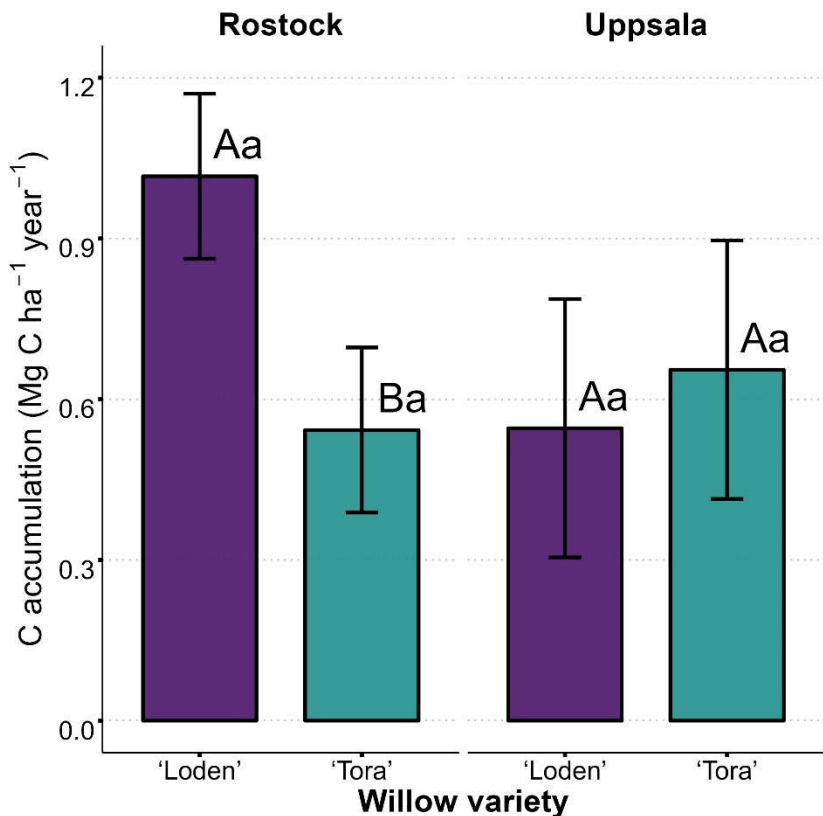


Figure 1. Soil C accumulation: estimated marginal means of topsoil (0-10 cm depth) C accumulation rates for the Salix varieties 'Loden' and 'Tora' grown in monoculture at sites Rostock and Uppsala. Site-specific bulk densities 1.3 and 1.4 g cm³ used in calculations of C stocks in Rostock and Uppsala respectively. Different uppercase letters indicate significant pairwise differences between willow varieties within each site (A-B; $p < 0.05$). Different lowercase letters indicate significant pairwise differences for each willow variety between sites (a-b; $p < 0.05$). Standard errors are shown as error bars.

The SOM chemical composition differed between 'Loden' and 'Tora' ($p = 0.036$) and between the mixture and 'Tora' ($p = 0.043$) along PC1 at the Rostock site (Figure 2). No significant differences between 'Loden', 'Loden': 'Tora' or 'Tora' were found along PC1 at the Uppsala site, or along PC2 at either site. Principal component 1 (PC1) explained 92.4% of the variation in the 9 compound classes from Py-FI mass spectra (Table 2) of soil samples under the willow varieties in monoculture and mixture and was strongly associated ($p < 0.001$) with all nine compound classes. Principal component 2 (PC2) explained 3.7% of the variation and correlated with compound classes lignin dimers (LDIM), lipids (LIPID) and free fatty acids (FATTY) ($p < 0.05$).

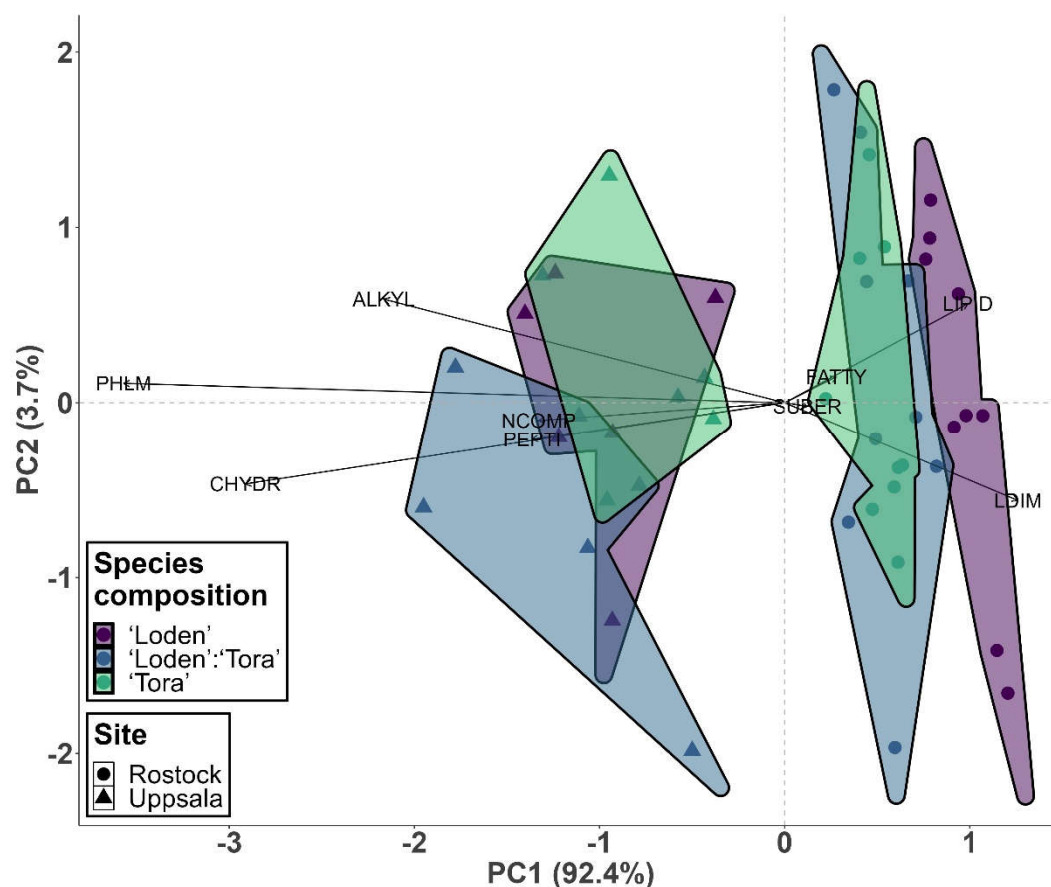


Figure 2. SOM chemical composition: principal components plot of the relative ion intensities (% TII) of nine compound classes; CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER, FATTY in Py-FI mass spectra (for abbreviations, see Table 2) in topsoil (0-10 cm depth) for the *Salix* varieties 'Loden' and 'Tora' grown in monoculture and mixture at sites Rostock and Uppsala.

The SOM under 'Tora' exhibited significantly greater abundances of carbohydrates (CHYDR; 1.0% TII, $p = 0.015$), phenols and lignin monomers (PHLM; 1.8% TII, $p = 0.018$), N-compounds (NCOMP; 0.5% TII, $p = 0.017$), and peptides (PEPTI; 0.6% TII, $p = 0.026$), but significantly lower abundances of suberin (SUBER; -0.2% TII, $p = 0.014$) and free fatty acids (FATTY; -0.6% TII, $p = 0.014$), compared to SOM under 'Loden' at the Rostock site (Table 5). Conversely, at Uppsala, the abundance of lignin dimers (LDIM; 1.7% TII, $p < 0.001$) was significantly higher in SOM under 'Tora' than 'Loden', while peptides (PEPTI; -0.8% TII, $p < 0.001$) and free fatty acids (FATTY; -0.1% TII, $p = 0.004$) were significantly lower under 'Tora' compared to 'Loden'. Additionally, comparing SOM chemical composition under each willow variety between sites, 'Loden' cultivated in Uppsala showed significantly higher abundances of carbohydrates (CHYDR; 5.5% TII, $p = 0.003$), phenols and lignin monomers (PHLM; 7.3% TII, $p = 0.001$), alkylaromatics (ALKYL; 4.5% TII, $p = 0.002$), N-compounds (NCOMP; 2.6% TII, $p = 0.002$), and peptides (PEPTI; 2.8% TII, $p < 0.001$) than 'Loden' cultivated in Rostock. In contrast, compound classes lignin dimers (LDIM; -2.6% TII, $p = 0.040$), lipids (LIPID; -2.0% TII, $p = 0.006$), suberin (SUBER; -0.31% TII, $p = 0.001$), and free fatty acids (FATTY; -0.71% TII, $p = 0.008$) were significantly lower under 'Loden' when cultivated in Uppsala compared to Rostock. For 'Tora', significant differences between sites included higher abundances of carbohydrates (CHYDR; 3.8% TII, $p = 0.010$), phenols and lignin monomers (PHLM; 4.5% TII, $p = 0.008$), alkylaromatics (ALKYL; 2.9% TII, $p = 0.009$), N-compounds (NCOMP; 1.5% TII, $p = 0.016$) and peptides (PEPTI; 1.4% TII, $p = 0.012$), in Uppsala compared to Rostock. Conversely, abundances of lipids (LIPID; -1.2% TII, $p = 0.037$) and suberin (SUBER; -0.11% TII, $p = 0.038$) were significantly lower in Uppsala compared to Rostock under 'Tora'.

Table 5. SOM chemical composition: estimated marginal means (\pm SE) of relative ion intensities (% TII) of nine compound classes; CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER, FATTY in Py-FI mass spectra (for abbreviations, see Table 2) in topsoil (0-10 cm depth) for the *Salix* varieties 'Loden' and 'Tora' grown in monoculture at sites Rostock and Uppsala. Different uppercase letters indicate significant pairwise differences between willow varieties within each site (A-B; $p < 0.05$). Different lowercase letters indicate significant pairwise differences for each willow variety between sites (a-b; $p < 0.05$).

Compound classes	Rostock		Uppsala	
	'Loden'	'Tora'	'Loden'	'Tora'
CHYDR	4.0 (0.19) ^{Bb}	5.0 (0.19) ^{Ab}	9.5 (0.82) ^{Aa}	8.8 (0.82) ^{Aa}
PHLM	9.9 (0.36) ^{Bb}	11.7 (0.36) ^{Ab}	17.2 (0.82) ^{Aa}	16.2 (0.82) ^{Aa}
LDIM	6.8 (0.64) ^{Aa}	7.1 (0.64) ^{Aa}	4.2 (0.59) ^{Bb}	5.9 (0.59) ^{Aa}
LIPID	10.3 (0.29) ^{Aa}	9.9 (0.29) ^{Aa}	8.4 (0.24) ^{Ab}	8.8 (0.24) ^{Ab}
ALKYL	13.3 (0.49) ^{Ab}	14.9 (0.49) ^{Ab}	17.8 (0.38) ^{Aa}	17.9 (0.38) ^{Aa}
NCOMP	1.6 (0.10) ^{Bb}	2.1 (0.10) ^{Ab}	4.1 (0.36) ^{Aa}	3.6 (0.36) ^{Aa}
PEPTI	3.6 (0.24) ^{Bb}	4.1 (0.24) ^{Ab}	6.4 (0.21) ^{Aa}	5.5 (0.21) ^{Ba}
SUBER	0.33 (0.03) ^{Aa}	0.16 (0.03) ^{Ba}	0.02 (0.01) ^{Ab}	0.05 (0.01) ^{Ab}
FATTY	0.87 (0.13) ^{Aa}	0.31 (0.13) ^{Ba}	0.17 (0.07) ^{Ab}	0.04 (0.07) ^{Ba}

CO₂ respiration, which was measured only at the Uppsala site, was similar across all variety compositions (Table 3). Chemical diversity (H') was significantly higher under 'Loden' compared to 'Tora' at the Rostock site ($p = 0.028$), with no difference at the Uppsala site (Table 4). The ratio of microbial- to plant-derived sugars (hexoses:pentoses), an indicator of microbial contribution to SOM, was significantly higher in SOM under 'Tora' compared to 'Loden' and the 'Loden':'Tora' mixture at the Uppsala site ($p = 0.021$ and $p = 0.026$, respectively). Total ion intensity (TII) was similar within sites, while total thermal stability was lower under 'Loden':'Tora' compared to 'Loden' ($p = 0.018$) only at the Uppsala site. Volatile matter (VM) was lower under 'Loden':'Tora' compared to both 'Loden' and 'Tora' at the Uppsala site ($p < 0.001$ and $p = 0.003$, respectively).

The thermal stability of compound classes lignin dimers (LDIM; $p = 0.047$), lipids (LIPID; $p = 0.030$), alkylaromatics (ALKYL; $p = 0.032$), suberin (SUBER; $p = 0.015$) and free fatty acids (FATTY; $p = 0.033$) were all significantly higher under 'Tora' than 'Loden' at the Rostock site (Table 6 and Figure 3). In contrast, at the Uppsala site, only suberin had a greater thermal stability under 'Tora' ($p < 0.001$), representing the sole compound class that consistently responded to variety identity across sites. Compound classes carbohydrates (CHYDR; $p = 0.004$), phenols and lignin monomers (PHLM; $p = 0.006$), alkylaromatics ($p = 0.022$), N-compounds (NCOMP; $p = 0.009$), and peptides (PEPTI; $p < 0.001$) were all significantly more stable under 'Loden' than 'Tora' at the Uppsala site (Table 6 and Figure 4). In terms of comparisons of SOM thermal stability under each willow variety between sites, 'Loden' cultivated in Uppsala showed significantly higher thermal stability of compound classes phenols and lignin monomers (PHLM; $p = 0.033$), lipids (LIPID; $p = 0.014$), alkylaromatics (ALKYL; $p = 0.038$) and N-compounds (NCOMP; $p = 0.022$) compared to 'Loden' cultivated in Rostock. For 'Tora', no significant differences in SOM thermal stability were found between the sites.

Table 6. SOM thermal stability: estimated marginal means (\pm SE) of thermal stability (ions volatilized > 400 °C / ions volatilized 50-650 °C) of nine compound classes; CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER, FATTY in Py-FI mass spectra (for abbreviations, see Table 2) in topsoil (0-10 cm depth) for the *Salix* varieties 'Loden' and 'Tora' grown in monoculture at sites Rostock and Uppsala. Different lowercase letters indicate significant pairwise differences for each willow variety between sites (a-b; $p < 0.05$).

Compound classes	Rostock		Uppsala	
	'Loden'	'Tora'	'Loden'	'Tora'
CHYDR	0.32 (0.08) ^{Aa}	0.36 (0.08) ^{Aa}	0.60 (0.08) ^{Aa}	0.51 (0.08) ^{Ba}
PHLM	0.58 (0.06) ^{Ab}	0.67 (0.06) ^{Aa}	0.82 (0.05) ^{Aa}	0.77 (0.05) ^{Ba}

LDIM	0.88 (0.02) ^{Ba}	0.93 (0.02) ^{Aa}	0.95 (0.02) ^{Aa}	0.97 (0.02) ^{Aa}
LIPID	0.67 (0.04) ^{Bb}	0.80 (0.04) ^{Aa}	0.90 (0.03) ^{Aa}	0.90 (0.03) ^{Aa}
ALKYL	0.70 (0.04) ^{Bb}	0.80 (0.04) ^{Aa}	0.88 (0.04) ^{Aa}	0.87 (0.04) ^{Ba}
NCOMP	0.46 (0.07) ^{Ab}	0.53 (0.07) ^{Aa}	0.80 (0.06) ^{Aa}	0.68 (0.06) ^{Ba}
PEPTI	0.48 (0.07) ^{Aa}	0.54 (0.07) ^{Aa}	0.72 (0.06) ^{Aa}	0.64 (0.06) ^{Ba}
SUBER	0.76 (0.06) ^{Ba}	0.90 (0.06) ^{Aa}	0.88 (0.05) ^{Ba}	0.97 (0.05) ^{Aa}
FATTY	0.02 (0.01) ^{Ba}	0.07 (0.01) ^{Aa}	0.10 (0.07) ^{Aa}	0.12 (0.07) ^{Aa}

At the Rostock site, the mean thermal stability of each compound class was as follows: lignin dimers > suberin > alkylaromatics > lipids > phenols and lignin monomers > peptides > N-compounds > carbohydrates > free fatty acids (Figure 3).

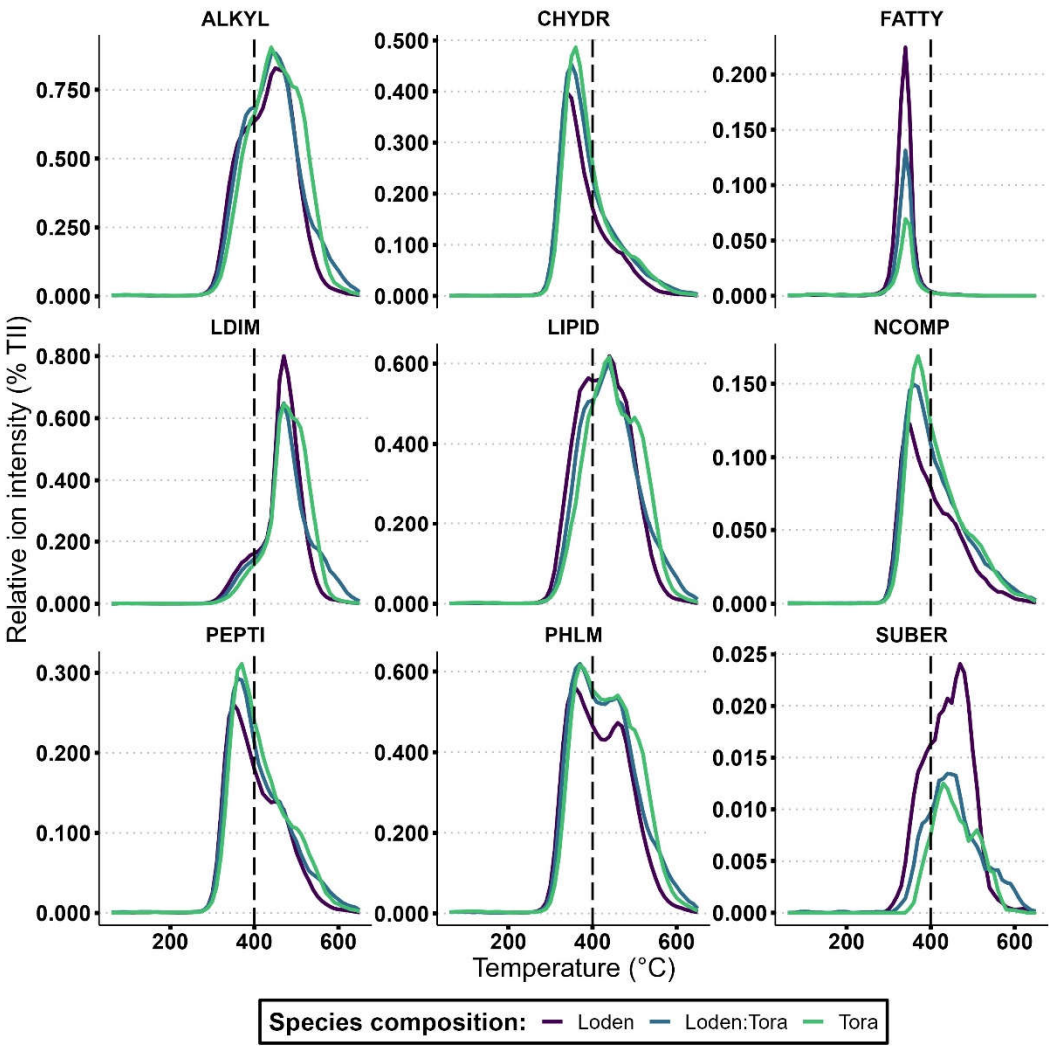


Figure 3. SOM thermal stability: thermograms showing ions volatilized under pyrolysis (0-650 °C) of nine compound classes; CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER, FATTY in Py-FI mass spectra (for abbreviations, see Table 2) in topsoil (0-10 cm depth) for the Salix varieties ‘Loden’ and ‘Tora’ grown in monoculture and mixture at the Rostock site. Higher values of relative ion intensity indicate a greater abundance of specific compounds in the pyrolyzed SOM. The temperature at which compounds volatilize reflects their thermal stability, with ions volatilized over 400 °C being regarded as stable and ions volatilized under 400 °C being regarded as labile.

At the Uppsala site, the mean thermal stability of each compound class was as follows: lignin dimers > lipids > suberin > alkylaromatics > phenols and lignin monomers > N-compounds > peptides > carbohydrates > free fatty acids (Figure 4).

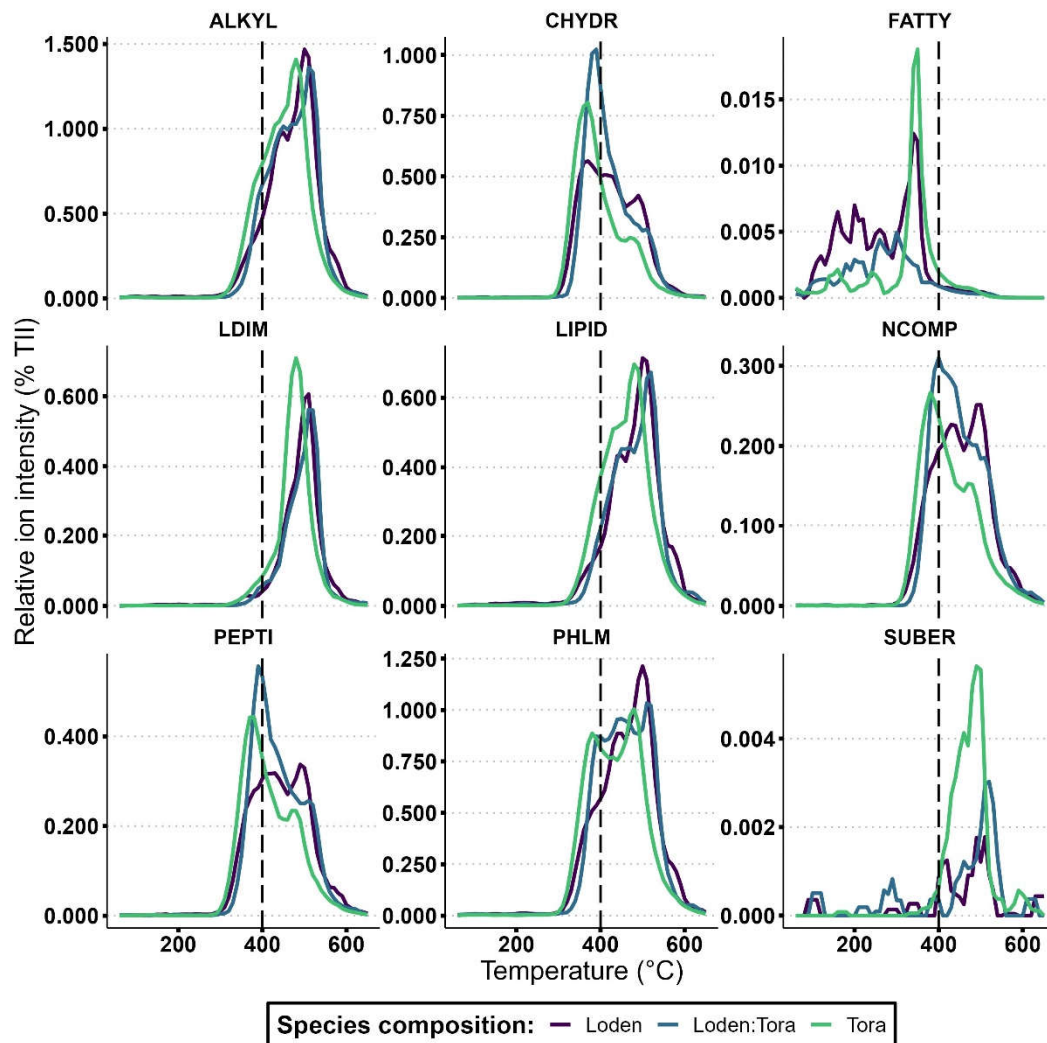


Figure 4. SOM thermal stability: thermograms showing ions volatilized under pyrolysis (0-650 °C) of nine compound classes; CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER, FATTY in Py-FI mass spectra (for abbreviations, see Table 2) in topsoil (0-10 cm depth) for the *Salix* varieties 'Loden' and 'Tora' grown in monoculture and mixture at the Uppsala site. Higher values of relative ion intensity indicate a greater abundance of specific compounds in the pyrolyzed SOM. The temperature at which compounds volatilize reflects their thermal stability, with ions volatilized over 400 °C being regarded as stable and ions volatilized under 400 °C being regarded as labile.

3.3. Effects of Variety Mixing

Expected values of yearly soil C accumulation in mixtures, as based on basal area weighted and averaged monoculture values, were similar to observed mixture values at either site (Figure 5). Expected and observed soil C accumulation in mixtures were 0.75 and 0.98 Mg C ha⁻¹ year⁻¹, respectively, at the Rostock site, and 0.61 and 0.16 Mg C ha⁻¹ year⁻¹, respectively, at the Uppsala site. At the Rostock site, observed mixture values were found to be significantly higher than at the Uppsala site ($p = 0.046$).

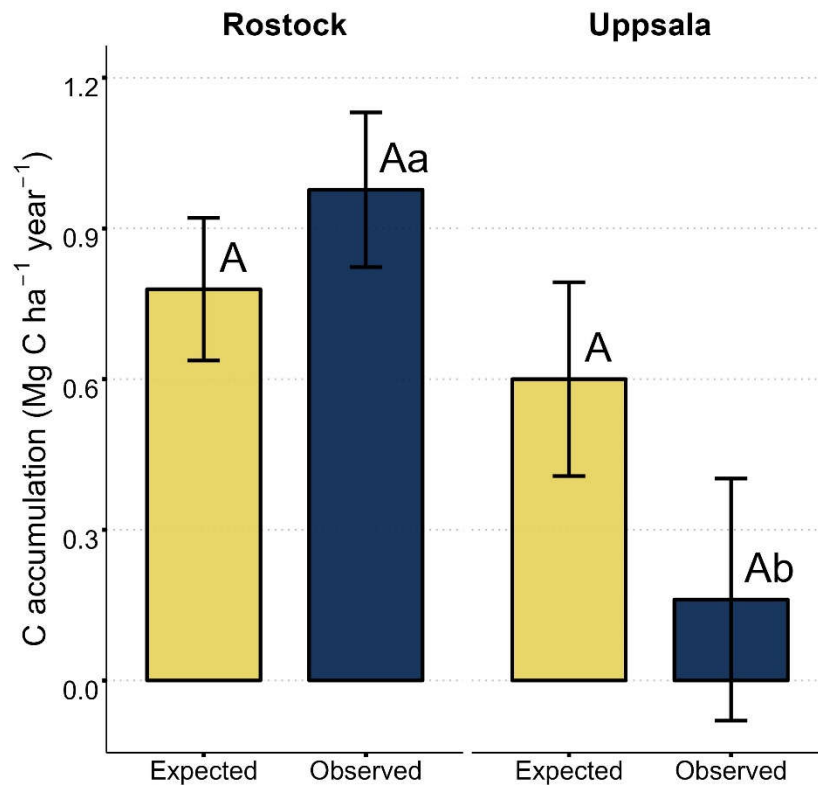


Figure 5. Soil C accumulation: estimated marginal means of expected and observed topsoil (0-10 cm depth) C accumulation rates at sites Rostock and Uppsala. Site-specific bulk densities 1.3 and 1.4 g cm³ used in calculations of C stocks in Rostock and Uppsala, respectively. Expected values represent the basal area weighted and averaged mean values of the *Salix* varieties 'Loden' and 'Tora' grown in monoculture and observed values represent the mean values of 'Loden' and 'Tora' grown in mixture (see Equation (3)). Different uppercase letters indicate significant pairwise differences between expected and observed mixture values within each site (A-B; $p < 0.05$). Different lowercase letters indicate significant pairwise differences for observed mixture values between sites (a-b; $p < 0.05$). Standard errors are shown as error bars.

Expected SOM chemical composition in mixtures, as based on basal area weighted and averaged monoculture values, were mostly similar to the observed mixture values (Table 7). Significant non-additive diversity effects were found for lignin dimers (LDIM) and peptides (PEPTI) at the Uppsala site, where lignin dimer abundance was reduced under mixtures (NDE = -18.3%, $p < 0.001$) compared to monocultures and peptide abundance was elevated under mixtures compared to monocultures (NDE = 18.2%, $p < 0.001$).

Table 7. SOM chemical composition: estimated marginal means (\pm SE) of expected and observed relative ion intensities (% TII) of nine compound classes: CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER and FATTY in Py-FI mass spectra (for abbreviations, see Table 2), as well as net diversity effect (NDE; Equation (3)) in topsoil (0-10 cm depth) at sites Rostock and Uppsala. Expected values represent the basal area weighted and averaged mean values of the Salix varieties 'Loden' and 'Tora' grown in monoculture and observed values represent the mean values of 'Loden' and 'Tora' grown in mixture (see Equation (3)). Asterisks indicate significant pairwise differences between expected and observed values within each site (*** = $p < 0.001$, ** = $p < 0.01$, * = $p < 0.05$, n.s. = not significant).

Compound classes	Rostock			Uppsala		
	Expected	Observed	NDE	Expected	Observed	NDE
CHYDR	4.6 (0.14)	5.0 (0.19)	9.2% n.s.	9.0 (0.62)	10.6 (0.82)	17.1% n.s.
PHLM	11.0 (0.26)	11.6 (0.36)	5.8% n.s.	16.5 (0.63)	17.4 (0.82)	5.7% n.s.
LDIM	7.0 (0.62)	6.6 (0.64)	-5.6% n.s.	5.4 (0.59)	4.4 (0.59)	-18.3% ***
LIPID	10.1 (0.23)	9.9 (0.29)	-1.9% n.s.	8.6 (0.20)	8.0 (0.24)	-7.2% n.s.
ALKYL	14.2 (0.35)	14.6 (0.49)	2.8% n.s.	17.8 (0.29)	17.6 (0.38)	-1.0% n.s.
NCOMP	1.9 (0.07)	2.1 (0.25)	10.2% n.s.	3.8 (0.27)	4.5 (0.23)	20.1% n.s.
PEPTI	3.9 (0.22)	4.1 (0.24)	4.0% n.s.	5.8 (0.20)	6.8 (0.21)	18.2% ***
SUBER	0.23 (0.03)	0.20 (0.03)	-14.8% n.s.	0.04 (0.01)	0.03 (0.01)	-18.0% n.s.
FATTY	0.55 (0.10)	0.52 (0.13)	-6.5% n.s.	0.08 (0.07)	0.13 (0.07)	59.4% n.s.

Comparable to SOM chemical composition, SOM thermal stability was mostly similar between expected and observed mixture values for most compound classes, though some significant non-additive mixture effects were found, but only at the Uppsala site (Table 8 and Figure 4). There, the stability of compound classes ALKYL (NDE = -0.9%, $p = 0.036$) and suberin (SUBER; NDE = -22.7%, $p < 0.001$) were both lower under mixtures compared to monocultures.

Table 8. SOM thermal stability: estimated marginal means (\pm SE) of expected and observed thermal stability (ions volatilized $> 400^\circ\text{C}$ / $> 0^\circ\text{C}$) of nine compound classes: CHYDR, PHLM, LDIM, LIPID, ALKYL, NCOMP, PEPTI, SUBER and FATTY in Py-FI mass spectra (for abbreviations, see Table 2), as well as the net diversity effect (NDE; Equation (3)) in topsoil (0-10 cm depth) at sites Rostock and Uppsala. Expected values represent the basal area weighted and averaged mean values of varieties 'Loden' and 'Tora' grown in monoculture and observed values represent the mean values of 'Loden' and 'Tora' grown in mixture (see Equation (3)). Asterisks indicate significant pairwise differences between expected and observed values within each site (*** = $p < 0.001$, ** = $p < 0.01$, * = $p < 0.05$, n.s. = not significant).

Compound classes	Rostock			Uppsala		
	Expected	Observed	NDE	Expected	Observed	NDE
CHYDR	0.34 (0.08)	0.36 (0.08)	3.3% n.s.	0.54 (0.08)	0.53 (0.08)	-1.0% n.s.
PHLM	0.63 (0.05)	0.63 (0.06)	-1.0% n.s.	0.78 (0.05)	0.78 (0.05)	0.3% n.s.
LDIM	0.91 (0.02)	0.91 (0.02)	-0.6% n.s.	0.96 (0.02)	0.95 (0.02)	-1.5% n.s.
LIPID	0.74 (0.04)	0.74 (0.04)	-0.7% n.s.	0.90 (0.03)	0.90 (0.03)	0.4% n.s.
ALKYL	0.76 (0.04)	0.74 (0.04)	-2.0% n.s.	0.87 (0.04)	0.86 (0.04)	-0.9% *
NCOMP	0.48 (0.06)	0.50 (0.07)	-0.2% n.s.	0.71 (0.06)	0.74 (0.06)	3.55% n.s.
PEPTI	0.52 (0.07)	0.50 (0.07)	-2.5% n.s.	0.67 (0.07)	0.67 (0.06)	0.4% n.s.
SUBER	0.84 (0.05)	0.83 (0.06)	-1.8% n.s.	0.99 (0.05)	0.76 (0.05)	-22.7% ***
FATTY	0.05 (0.01)	0.04 (0.01)	-19.8% n.s.	0.12 (0.05)	0.15 (0.07)	30.8% n.s.

4. Discussion

4.1. Effects of Site-Specific Environmental Conditions

Our study used two experimental sites with distinct climatic and edaphic circumstances, which provided a platform to investigate how local conditions modulate the influence of willows on SOM quality. Our results showed that the interaction between plants and the investigated soil properties is strongly influenced by local site conditions, where soil C accumulation, SOM chemical composition and thermal stability responded differently to both variety identity and mixture across sites (confirming hypothesis 3). Temperature, moisture and especially clay content, which differed between the two study sites, are widely documented as strong modifiers of decomposition rates [23,81], and microbial dynamics [82–84]. Therefore, we expected the lower temperatures and precipitation levels as well as higher clay content in Uppsala to contribute to lower SOM decomposition rates compared to Rostock. In the present study, this is partly supported by the lower ratio of microbial- to plant-derived sugars (hexoses:pentoses) found in Uppsala compared to Rostock, indicating lower microbial contribution to SOM [70,85]. Additional supporting evidence for lower decomposition rates in Uppsala compared to Rostock comes from a previous study on the same willow SRCs [60], showing slower leaf litter decomposition rates for monocultures and higher fractions of remaining N (immobilization of N) in mixtures in Uppsala compared to Rostock, though with only minor effects of climate. However, the highest accumulation of soil C was found under 'Loden' in Rostock, where we expected a higher decomposition rate compared to Uppsala. This discrepancy between likely outcomes based on literature evaluations and results may be explained by the generally greater biomass production at the Rostock site [64], leading to a greater litter input to the soil.

4.2. Effects of Willow Variety

In the present study, we demonstrated that the influence of willow SRC on soil C accumulation and SOM chemistry strongly depended on willow variety (confirming hypothesis 1). After seven years of growth, the net average annual C accumulation in the topsoil (0-10 cm depth) was consistently positive across all experimental plots, with C accumulation rates ranging from 0.06 to 1.29 Mg C ha⁻¹ year⁻¹. These values were comparable to mean soil C accumulation values documented under willow SRC for 'Loden' (0.73 Mg C ha⁻¹ year⁻¹) and 'Tora' (0.65 Mg C ha⁻¹ year⁻¹) near the Uppsala site over a 17 year period in [51]. In the site with climatically more favorable growth conditions (Rostock), 'Loden' had a higher C accumulation rate compared to 'Tora', whilst both varieties accumulated similar soil C amounts at the site with climatically less favorable growth conditions (Uppsala). A greater C accumulation under 'Loden' could be due to a higher fine root biomass production compared to 'Tora', as reported by [57] at the Rostock site. Furthermore, at the Rostock site, 'Loden' also had higher abundances of suberin compared to 'Tora', a compound primarily derived from roots [86]. This suggests a higher fine root biomass production under 'Loden' at the Rostock site, thus in line with the findings of [57]. Interestingly, the difference in C accumulation between 'Loden' and 'Tora' in Rostock was similar to what [51] reported in a site near Uppsala after 17 years of growth. It is therefore plausible that we in the future (e.g., after an additional 10 years) will see a significantly greater soil C accumulation under 'Loden' than 'Tora' also in the site with less favorable growth conditions (Uppsala). The difference in C accumulation between 'Loden' and 'Tora' in Rostock could also depend on differences in SOM chemistry, or results in different stability of the accumulated SOM. At the Rostock site, we found more chemically diverse SOM under 'Loden' compared to 'Tora', which may have contributed to a decreased decomposition and a subsequently promoted C accumulation for 'Loden' [4,87]. The relative enrichment of monomeric lignin building blocks under 'Tora' at the Rostock site, which can be explained by an advanced lignin decomposition [15], also supports enhanced decomposition in 'Tora' plots. Further, 'Tora' plots at the Rostock site displayed higher overall thermal stabilities of compound classes, suggesting higher decomposition rates as the more labile fractions of plant litter SOM would be decomposed first [11], leaving more recalcitrant molecules behind. A final potential explanation to the differences in C

accumulation between 'Loden' and 'Tora', in addition to differential biomass production and decomposition rates, can depend on the interaction between mycorrhizal and saprotrophic fungi. Since 'Loden' is more dependent on the ectomycorrhizal association compared to 'Tora' [57,63] the saprotrophic activity under 'Loden' may have been suppressed as per the Gadgil effect [88]. The Gadgil effect implies suppression of saprotrophic activity in the presence of increased ectomycorrhizal colonization due to competition between the two functional groups and assimilate transfer from the host plant.

4.3. Effects of Variety Mixing

In the present study, we found no effect of variety mixing on soil C accumulation (confirming part of hypothesis 2), which is consistent with the findings of [89–91]. Since previous investigations of mixed-species willow plantations with 'Loden' and 'Tora' have found no indication of aboveground over-yielding [64,92], which is considered one of the main mechanisms of increased soil C sequestration in mixtures [25,93], this may explain the lack of response. However, even without increased plant productivity, 'Loden':'Tora' mixtures have been shown to affect other belowground processes such as soil phosphorus (P) cycling [57] e.g., through promoting alkaline phosphatase activity in P-deficient soils in mixtures compared to monocultures [92], and to increase endophytic root colonization [57], when compared to their monoculture counterparts. We detected some significant effects of variety mixing on SOM chemical composition and thermal stability (confirming the second part of hypothesis 2). At the Uppsala site, the detected effects of variety mixing on SOM chemical composition were found in the compound classes of lignin dimers and peptides, which revealed significantly higher and lower abundances, respectively, in variety mixtures compared to equivalent monocultures. Elevated levels of (lignin) decomposition in mixtures as a response to litter diversity is a possible explanation for these findings [94], which might be based on the higher general decomposition activity as revealed for the P cycling [57]. Since lignin is a rather stable fraction of SOM, lower lignin levels in mixtures could reduce SOM stability and decrease the long-term C sequestration in mixtures. Even so, the absence of effects of variety mixing on soil C accumulation and only some effects on SOM chemistry suggests that mixed willow cultivation does not greatly impair SOM quality or climate change mitigation potential compared to equivalent monoculture plantations at the same location. However, an increased decomposition of lignin in the mixtures in Uppsala could also indicate a decreased SOM formation. Finally, soil respiration, which is largely based on microbial activity in the soil, can reflect increased decomposition rates when respiration rates are higher. However, we did not find any difference in soil respiration between the treatments in the present study.

5. Conclusions

Our results highlight the complexity of plant-soil interactions and the challenges they present during interpretation. However, despite the complex nature of our results, our study revealed that willow variety or species identity could have a significant impact on soil C accumulation, SOM chemical composition and thermal stability. We found no evidence for non-additive effects of willow variety mixtures on soil C accumulation, but some non-additive effects on SOM chemical composition and thermal stability. Significant interaction between willow variety and site conditions have important implications for the management of willow SRC plantations. First, variety or species mixtures can be used in multifunctional plantations without greatly compromising soil C accumulation or SOM quality, as they mainly exhibit values similar to their monoculture counterparts. Second, the selection of willow varieties or species for willow SRC plantations need to consider the context-dependency of their expression under specific site conditions.

Supplementary Materials: The following supporting information can be downloaded at the website of this paper posted on Preprints.org, Figure S1: PCA using Py-FI mass signals (m/z) instead of compound classes; Table S1: Basal area weights used for calculating net diversity effects (NDE); Table S2: Soil chemical characteristics: results of two-way ANOVAs for soil C:N ratios nutrient concentrations and soil CO₂ respiration; Table S3: Py-FIMS parameters: results of two-way ANOVAs for various SOM properties; Table S4: Soil C

accumulation: results of a two-way ANOVA of topsoil C accumulation rates; Table S5: Principal component regression: results of two-way ANOVAs for principal components (PC1 and PC2) derived from the relative ion intensity (% TII) of nine compound classes in topsoil; Table S6: SOM chemical composition: results of two-way ANOVAs for the relative ion intensity (% TII) of nine compound classes in Py-FI mass spectra in topsoil; Table S7: SOM thermal stability: results of two-way ANOVAs for the thermal stability of nine compound classes in Py-FI mass spectra in topsoil; Table S8: Py-FI mass spectral data of soil samples from Uppsala and Rostock; Table S9: C stocks and accumulation rates as well as P, Mg and K concentrations in topsoil.

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