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1 Abstract

During the last centuries, land use in Europe intensified, which has led to a drastic decrease in the cover of semi-natural grasslands. In Sweden, much of the lost grasslands was turned into forest. This study investigated if species typical of managed grasslands could be found in coniferous production forests more than 80 years after grassland management ceased. Species and trait composition for plants was investigated in two types of forest differing in land use history (meadow in the 1870s or continuous coniferous forest), and in reference grasslands. The average plant species richness as well as the richness of grassland indicator species were 30% higher in forests with a history as meadow compared to in forests with a history as forest, hence clear signs of historical grassland management in today's forests. Compared with forests with continuous coniferous history, vegetation in forests with a meadow history tended to be more similar to reference grassland regarding both plant species and especially plant trait composition. The study provides proof of remnant grassland populations in coniferous production as the source for the biodiversity of clearcuts, rather than seed dispersal or seed bank survival. The result highlights the importance of land use for biodiversity of clearcuts, and points to

the potential value of forests with a history of meadow in grassland conservation and restoration.

2 Introduction

It is well-known that species indicative of previous land use can reappear after a disturbance or when management changes (e.g. Vellend et al. 2006, Hermy & Verheyen 2007, Szabó 2010, Munteanu et al. 2015). Such re-appearances contribute to population persistence and genetics, and they can also be considered as a natural or cultural legacy of the land (Foster et. al 2003, Eriksson & Cousins 2014). Furthermore, species' re-appearance is a key element of ecological restoration. Two obvious key questions follow regarding reappearances: what is the source for observed recolonisation? And for how long can we expect such recolonization to occur under a land use hostile to the species in question?

A major disturbance affecting a substantial part of forested landscapes is clearcutting, causing drastic changes in vegetation (Bergstedt & Milberg 2001, Bergstedt et al. 2008, Cesoniene et al. 2018). Evidence is accumulating that reappearance of plants can happen after at least a forest rotation (i.e. 80-120 yrs), i.e. in conjunction with clearcutting (Ibbe et al. 2011, Risberg & Granström 2012, Jonason et al. 2014). But within the forest/clearcut system, the source for observed recolonisation remains unclear. Jonason et al. (2014) addressed two hypotheses. Firstly, the seed bank hypothesis, i.e. that species with persistent seed bank should be more prevalent on clearcuts with a land use legacy as meadow compared with clearcuts in places with continuous coniferous forest cover. Secondly, the dispersal hypothesis, i.e. species with adaptation for dispersal would be more prevalent on clearcuts with a history of meadow compared with continuous forest cover. Jonason et al. (2014) could find no support for any of these, and conclude that a third hypothesis, that species persist as growing specimens through the forestry rotation seemed a more likely explanation (Jonason et al. 2014). Here, we call this the perseverance hypothesis. The fact that plants can survive in conditions that are relatively hostile is a well-known phenomenon (e.g. Eriksson 1996, Bond & Midgley 2001), and is contributing to situations of "extinction debt" (Helm et al. 2006, Kuusaari et al. 2009), i.e. a time-lag in loss of species in a community. From the point of view of population dynamics, perseverance would mean having a growth rate (k) below 1.

In the present contribution, we wanted to evaluate the perseverance hypothesis, essential to understand, and exploit, the naturally occurring potential in forestry and ecological restoration. We did this by vegetation inventories in fully mature forests, but with different land use history (forest or meadow in 1870s), destined for clearcut. We also wanted to evaluate if ecological filtering occurred among plant traits during the vegetation change in the chronosequence from open grassland to mature forest. This transition spans more than 100 yrs in our study system and involves elimination of annual mowing or domestic grazing as well as increased shade from trees. Plant traits has proven to be a useful tool to understand plant responses to environmental change (Lavorel & Garnier 2002), and we expected ecological filtering, i.e. a convergence in traits relevant for the secondary succession from grassland to forests (Kahmen & Poschlod 2004, Dölle et al. 2008), assessed by shifts in SD of traits. Our hypothesis about ecological filtering predicts that SD of plant traits would be higher in grasslands than in forests with a history as grassland (meadow). The actual changes in trait averages, however, were not of interest to us in the current contribution. We assessed both traits clearly expected to be affected (plant height, grazing/mowing tolerance, Ellenberg light value) and traits less apparently affected: lifeform, Ellenberg moisture, Ellenberg pH, Ellenberg nitrogen and

seed traits (diaspore mass, terminal velocity of diaspore [maximum speed of falling seed], epizoochory, seed bank).

3 Material & methods

3.1 Study sites

The study was performed in southern Sweden in the province of Östergötland (Figure 1), which is in the hemiboreal zone of Scandinavia (Ahti et al. 1968). Forty sites were investigated, 12 semi-natural grasslands and 28 mature forests (which correspond to approximately 80-120 years), planned for clear-cutting within three years. The forests were evenly divided between forests with a management history as meadow or as coniferous forest according to land use maps from 1870s. All forests were coniferous production forests with either Norway spruce (*Picea abies*) or Scots pine (*Pinus sylvestris*) as the dominating tree species. The size of the forested areas varied between 2.0 and 7.5 ha, and grasslands between 2.6 and 6.3 ha.

Species-rich semi-natural grasslands were identified using a geographical database managed by the Swedish Board of Agriculture (TUVA database; <u>www.sjv.se/tuva</u>). Such grasslands are generally believed to have a long continuity or management as meadow and/or grazing (e.g. Eriksson et al. 2002, Dahlström et al. 2006, Fredh et al. 2012, Bergstedt et al. 2017).

Forest data were provided by the Swedish Forest Agency to which landowners are obligated to report planned clear-cutting (Skogsdataportalen; <u>http://skogsdataportalen.skogsstyrelsen.se/Skogsdataportalen/</u>). To find suitable forests, historical land-use maps from the 1870s (Häradsekonomiska kartan) were compared with data on forests that were planned for clear-cutting. The old land-use maps contain information about agricultural land, coniferous forests, deciduous forest, meadows, wetlands, roads and more. Although surveyed and printed in the 1870s, part of the information was based on older cadastral maps (Jansson 1993, Runborg 1994). The sampled forests and old meadows did not overlap perfectly and for a forest to be categorized as history as meadow at least 25% of the area had to be meadow during 1870; the whole forest was treated as one unit in the analyses. The reasons for this are (i) the poor micro-spatial accuracy of the old land-use maps, (ii) errors introduced during digitalization e.g. due to rectification, (iii) that borders have shifted, and (iv) that the unit targeted by forestry operations is a forest stand irrespective of previous land use. Furthermore, (v) it is unclear how distinct old land use borders were because even if meadows were always fenced, there were some amount of change by previous meadows falling out of use and new areas taken in. Finally, (vi) grazing is likely to have occurred near meadows which suggest a grazed and relatively open forest near meadows.

Exactly when a former meadow was transformed to production forest is not known. Each area has, however, at least hosted one generation of coniferous forest, which in Östergötland corresponds to approximately 80 years. For the areas with a history as coniferous forest we cannot rule out that they were used as meadow prior to 1870 although we deem it as unlikely since the large-scale transformation from meadow to forest occurred later, during the 20th century (Ihse 1995, Eriksson et al. 2002). To reduce any possible influence of nearby semi-natural grasslands, all forests selected were at least 300 m from any present-day grassland. To assure that forests selected carried separate populations, the distance between them was also at least 300 m, which is a distance beyond the average dispersal distance for plant species (Thomson et al. 2011).

3.2 Plant survey

The species richness of herbaceous plants was investigated once at each site between early August and late October 2014. Due to phenological variation within the season some species may be underrepresented in the survey (Bergfur et al. 2004), for example species with an early growth peak or senescence of leaves, but this was not expected to bias the result. In each site, one-hundred circular sample plots (radius 1 m) were evenly distributed along transects 25 meters apart. Within each sample plot, plants present were recorded and plant frequency was calculated as the number of sample plots in which a species was presence. Plants that could not be identified with certainty to species level, mainly species belonging to the family Poaceae and the genus *Carex*, were excluded from the data. *Melampyrum pratense* and *M. sylvaticum* were difficult to separate in the field and were therefore grouped together.

To classify species as indicators for semi-natural grasslands, the indicator systems of Ekstam & Forshed (1992) and Bertilsson & Paltto (2003) were used. Bertilsson and Paltto (2003) is a regional indicator system created to investigate grazing management quality in semi-natural grasslands in the province of Västergötland in southern Sweden. The book by Ekstam and Forshed (1992) does not provide a clear classification of species as indicators for grasslands. The aim of the book is instead to provide a tool for answering questions about grazing, mowing and vegetation development in semi-natural grasslands in Sweden. The book contains tables where species are given several indicator values, for example successional category, light demand, nitrogen demand, species as indicators for semi-natural grasslands, we used the successional category, an indicator that puts species into one of four successional categories (A – D) which is based on when species is lost in an abandoned grassland. (A) concerns species that increase or have a relatively unchanged abundance during

the first and second year after abandonment but decreases or disappear after three to five years; (B) is species that increase or have a relatively unchanged abundance during the first five years after abandonment but decrease or disappear after ten to fifteen years; (C) Species that increase or have a relatively unchanged abundance during the first fifteen years after abandonment but decrease or disappear after twenty-five to thirty-five years; (D) is species that have their strongest populations in a forested phase in the succession but increase in abundance directly after abandonment. We considered species in (A) and (B) as indicators for semi-natural grasslands. Final classification means a species has been scored as an indicator in at least one of the two indicator systems (Appendix 1).

3.3 Habitat factors

At every third plant sample plot, habitat factors were investigated within an area of 100 m^2 (circle with radius 5.64 m). The species and abundance of living trees and stumps >10 cm in diameter were recorded. The size of the living trees was measured at breast height and recorded in diameter classes (10-15 cm, 15-20, 20-25, etc). These values were converted to basal area per each site. The percent cover of bare rock, residues (from previous thinning or preparing for the clearcut), and exposed mineral soil within the circles were estimated visually. All these factors were investigated to ensure that there was no systematic difference between the two types of forest sites.

3.4 Plant traits

To investigate to what extent different plant traits were associated with a certain land-use history, several traits related to dispersal, persistence and habitat requirements were selected for analysis (Eriksson 1996, Fischer & Stöcklin 1997, Stöcklin & Fischer 1999, Johansson et al. 2011). A total of 11 traits were chosen for the analysis (Table 2) and data were taken from the LEDA Trait base (Kleyer et al. 2008), the Dispersal and Diaspore Database (Hintze et al. 2013) and a database on traits of plants of southern Sweden (Tyler & Olsson 2013). In these databases, trait values are missing for some species, therefore, all traits could not be assigned to all species. However, at least 65% (range 65-99%) of each trait could be assigned of all species.

3.5 Statistical analyses

To investigate if there were any systematic differences between forests with a history as forest or as meadow, mean values and corresponding 95% confidence intervals were calculated for several habitat factors (Table 1). To describe an individual species' association with forest type, an odds ratio (OR) was calculated (Rita & Komonen 2008): OR = (a / b) / (c / d), where "a" is the number of plots in continuous coniferous forests with species *i* occurring and "b" the total number of such plots; "c" is the number of plots in forest with a history as meadow where species *i* occurs and "d" the total number of such plots. The natural logarithm of the OR and the corresponding 95% confidence interval was then calculated, ln (OR) $\pm 1.96 * \sqrt{(1/a + 1/b + 1/c + 1/d)}$. A negative ln (OR) indicates association with forest with a history as meadow compared to as forest. Only species present in at least two forest sites were included in this analysis (Appendix 2).

To further illustrate the dissimilarities in vegetation composition, data from the two forest types as well as of reference grasslands, were subjected to Principal Component Analysis (PCA, using the CANOCO 5 software: ter Braak & Šmilauer 2012). These analyses were based on the abundance of 168 plant species at 40 sites (12 grasslands, 28 forests with different history). To investigate if certain plant species traits are associated with a certain site type, trait values per site were calculated as a community-weighted mean (CWM). CWM is the mean trait value of all species present in the community (site)

weighted by their relative abundances (in our case the frequency of occurrence in the 100 sample plots). Before the CWM was calculated, the species' frequency data were log-transformed in order to reduce the influence of dominating species. The traits used in the analyses consisted of several different variable types; continuous, rank, categorical and binary (Table 2). However, when calculating the CWM all traits were treated as continuous variables.

For each trait, the average value was calculated per forest or grassland. The mean and SD over grasslands, and each of the two forest types were calculated. In addition, corresponding means were calculated weighing each species by its frequency in sample plots. We used the SD and weighted SD to test the hypothesis about ecological filtering.

4 Results

In total, 168 herbaceous plant species were found during the survey in forests and grasslands (133 in grasslands; 134 in forest with a history as meadow; 90 in forests with a history as forest) (Appendix 1). Out of these, 48 species were classified as grassland indicator species (46 in Grasslands; 27 in forest with a history as meadow; 19 in forests with a history as forest). The average species richness was 29.6% higher in forests with a history as meadow compared to in forests with a history as forest (Figure 2a) and the richness of grassland indicator species was on average 30.1% higher in forests with a history as meadow (Figure 2b).

Most species were more frequent in forests with a history as meadow compared to forests with a history as forest (Figure 3). A total of 107 species were included in species-wise analyses and 53 of them showed distinct association (Cl_{95%} did not overlap zero) towards forests with a history as meadow (Figure 3), compared to 14 for forest towards a history as forest. Out of 23 grassland indicator species occurring in the forest plots, 10 showed a distinct preference

towards forest with a history as meadow and 2 showed preference for forests with a history as forest.

Land use history affected both the species (Figure 4) and trait composition (Figure 5) in forests. A meadow-history of forests showed a tendency to be more similar to grasslands regarding both species (Figure 4) and trait composition (Figure 5) compared to areas with a forest history. When analysing community-weighted traits, grasslands were distinctly different from forests mainly in Ellenberg light, Grazing tolerance and Seed bank persistence (Figure 5a). In contrast, only one trait was larger in coniferous forest continuum: terminal velocity (Figure 5a).

Our hypothesis that ecological filtering, executed by reduced grazing and/or light, is important when turning from grassland to forest, predicts that SD of plant traits would be higher in grasslands than in forests with a history as grassland (meadow). There were, however, no significant differences in the SD, nor the weighted SD, for any of the 11 traits evaluated (Table 3). In two cases there was a tendency for a difference, but both involved grasslands having the lower SD, contrary to the hypothesis.

5 Discussion

5.1 Land use legacy and perseverance

The results of this study clearly demonstrated that land use history can influence present-day plant species and trait composition in hemiboreal forests of Sweden with total species richness and richness of grassland indicator species being higher, on average, in forests with a history as meadow compared to in forests with a history as forest. Other studies have also shown that historical land use affect present day flora, for example in agricultural landscapes (Gustavsson et al. 2007) and clear-cuts (Jonason et al. 2014, 2016). Effects have also been

shown for other organisms such as butterflies (Ibbe et al. 2011, Blixt et al. 2015), also one study showed that watershed land use in the 1950 affected present day diversity of stream invertebrates and fishes (Harding et al. 1998). Historical land use can also affect the structure and function of whole ecosystems and it has been shown to be important in different biomes and habitats (Foster et al. 2003). The effects of land use can last for centuries (Dahlström et al. 2006, Fortuny et al. 2014) or in some cases even for millennia, as studies have shown that the farming intensity during the period AD 50-250 influenced species richness and plant composition today (Dupouey et al. 2002). The current study showed that remnant population of typical grassland species survive at least 80 years after that grassland management has changed to coniferous production forest. As mentioned above, Jonason et al. (2014) proposed three hypotheses for the source of these grassland species' occurrence on clearcuts: seed bank in soil, dispersal from nearby grasslands or perseverance in the shade of forest. They could find no support of the first two, while the present study provided clear support for the perseverance hypothesis, i.e. populations of grassland species persist in the shade of coniferous forest as remnant populations (Eriksson 1996, Dahlström et al. 2006). Hence, certain grassland species can persist in a changed habitat following the transitions of land use (meadow to forest to clear-cut) that occurred in southern Sweden during the last century. Or put another way, grassland species in forest that used to be grassland are subject to extinction debt (Kuussaari et al. 2009) but instalments are few and spaced in time.

Even if there is strong support for the perseverance hypothesis, this does not rule out that seed dispersal and/or seed bank may play some role too. For example, seed dispersal within a forest might help population survival by redistributing seeds from sun-light areas like verges or glades to darker areas. Furthermore, seed bank survival might help boost population growth after a clear cut.

5.2 Traits

During the transition from grassland to dense coniferous forest, drastic changes occur in response to reduced light and reduced grazing. This was clearly reflected in the community-weighted trait estimates for Ellenberg light and grazing tolerance, that both decreased. Grazing is an obvious factor that maintains grassland richness and composition (Dupré & Diekmann 2001, Öckinger et al. 2006, Milberg et al. 2017) and it is well-established that shading is detrimental to many grassland species (Einarsson & Milberg 1999, Skornik et al. 2008). Therefore, we had assumed that ecological filtering would reduce the variation in Ellenberg light value due to increased shading during the succession from grassland to mature forest, i.e. that the standard deviation (SD) of this trait would be smaller in forest. At the same time, the elimination of grazing would increase variation in plant height and grazing/mowing tolerance. We found no support for these assumptions in our data, nor for change in SD in the other traits evaluated. It remains open to speculation whether this is due to design weakness (space-for-time design), or to a faulty hypothesis.

5.3 Implications for conservation and restoration

The presence of grassland plant species in forest, as a legacy of earlier land use, indicate a conservation potential. Some of these plants have become rare today and are in need of an increase in suitable habitat to ensure long-term survival (Nilsson et al. 2013, Hoekstra et al. 2005). Studies have shown that restoration management by grazing after clear-cutting on sites with grassland history have a positive effect on the grassland flora (Piqueray et al. 2015) and that grassland restoration is possible decades after abandonment (Skórka 2007). Therefore, there is a potential for grassland restoration in forests with meadow history by

introducing grazing or mowing after clearcut (Van Uytvanck & Verheyen 2014). Old land use information may also be valuable when selecting sites for restauration or creation of woodland pastures (Roellig et al. 2016).

Remnant grassland species that flourish after a clearcut can provide nectar and other resources for species that we normally associate with grasslands (Viljur & Teder 2018, Ohwaki et al. 2018, Bergman et al. 2018). A practical issue is then how targeted populations of plants can be boosted. This can be done by postponing reforestation, which should give remnant populations more time to expand on the area and give new species confined to open areas a chance to establish. To maintain favourable conditions for grassland species during afforestation, some parts of the forests could be kept open and secondary succession allowed to proceed in the absence of planted trees. It is also likely that planting deciduous trees will prolong the time period that clearcuts are favourable for grassland species.

Studies have shown that plant species richness is higher in semi-natural grasslands (Söderström et al. 2001, Öckinger et al. 2012), road verges, midfield islets (Lindborg et al. 2014) and grasslands on former arable fields (Cousins & Aggemyr 2008) in forested landscapes compared to agricultural landscapes. This "supportive" influence of forests suggests that grassland populations might not be as fragmented as generally thought.

The current study points to the potential for conservation of grassland species in forests with history as meadow, either by restoration, modified re-forestation methods, or retention forestry with focus on the grassland species. With proper management there is potential to strengthen typical grassland populations in the landscape, both by increasing the chance of remnant populations surviving and by enhancing the connectivity of suitable habitats for the species in the landscape.

5.4 Conclusions

This study shows that grassland species might be more common in the landscape than assumed due to remnant grassland populations in coniferous production forests with a history of grassland management. Furthermore, this study highlights the use of such forests in grassland conservation and restoration. Restoration by grazing animals after clear-cutting has been shown to be successful and favour a lot of grassland plant species (Piqueray et al. 2015). Conservation aims can also be achieved by postponing afforestation, mowing or grazing a time after harvest and planting deciduous trees instead of coniferous.

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Figure legends

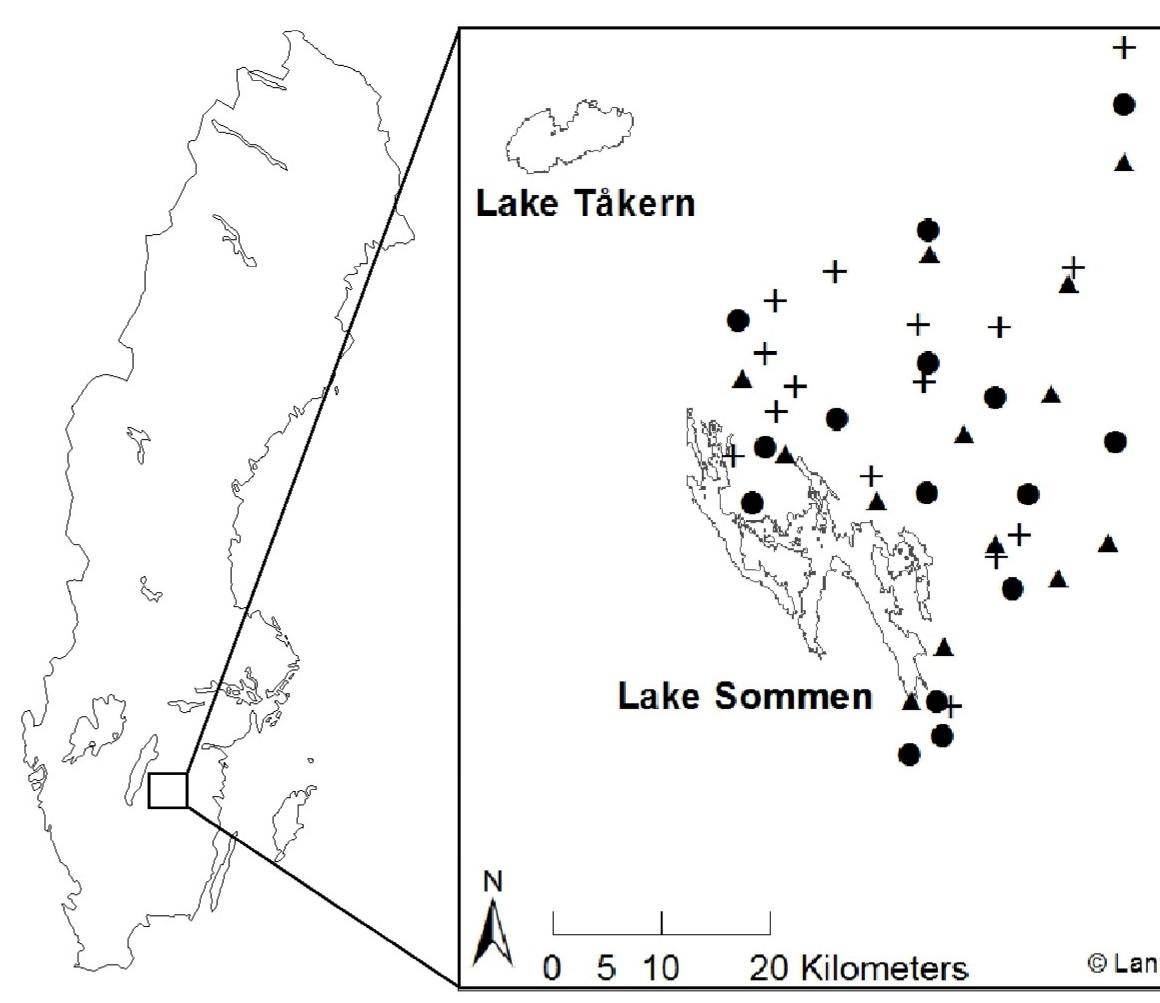
Figure 1. Map over Sweden (left) and the study area in the southern part of the province Östergötland (right).

Figure 2. Average plant species richness, with $Cl_{95\%}$, of forests with a history as forest (n=14) and as meadow (n=14) in the 1800s for (a) all species combined and for (b) species classified as grassland indicator species. The reference grassland sites had on average 74.2 species ($CI_{95\%}$ 68.6; 79.9) and 27.4 grassland indicator species (25.2; 29.6).

Figure 3. Odds ratio (natural logarithm) with CI_{95%} (whiskers) for species in forests differing in land use history. Positive values of a species indicate preference towards forests with history as forest and negative values indicates a preference towards forest with history as meadow. Size of the black symbol is proportional to the total frequency of a species. Species labelled with "*" are grassland indicator species.

Figure 4. PCA of species composition data from forest with different land use history (meadow or coniferous forest in the 1870s) and reference grasslands (12). Eigenvalues of PC1 and PC2 were 51.2 and 9.2%, respectively. Arrows indicate species.

Figure 5. PCA of community-weighted trait data from forest with different land use history (meadows or coniferous forest in the 1870s) and reference grasslands. Eigenvalues of PC1 and PC2 were 50.3 and 31.4%, respectively.

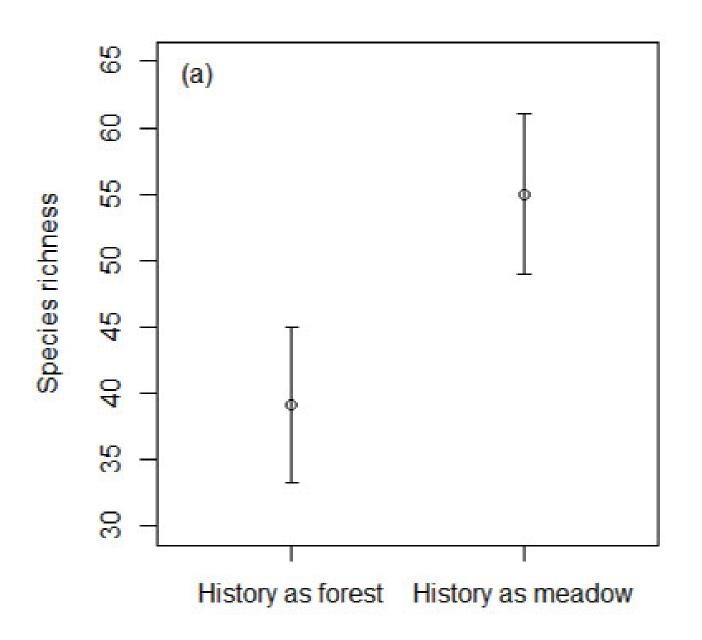


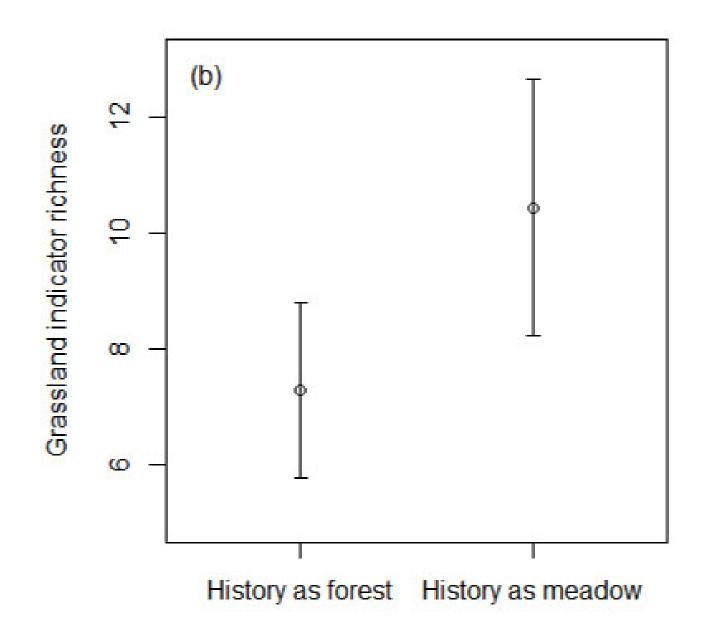
History as meadow

History as forest

Grasslands

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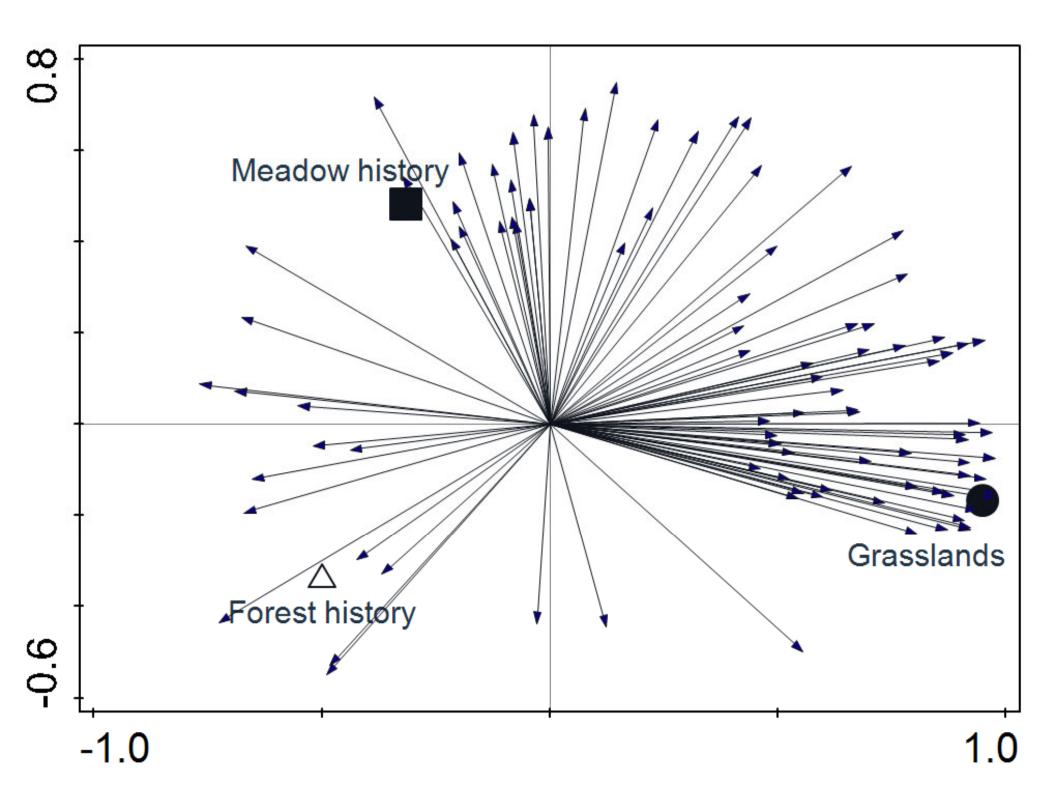


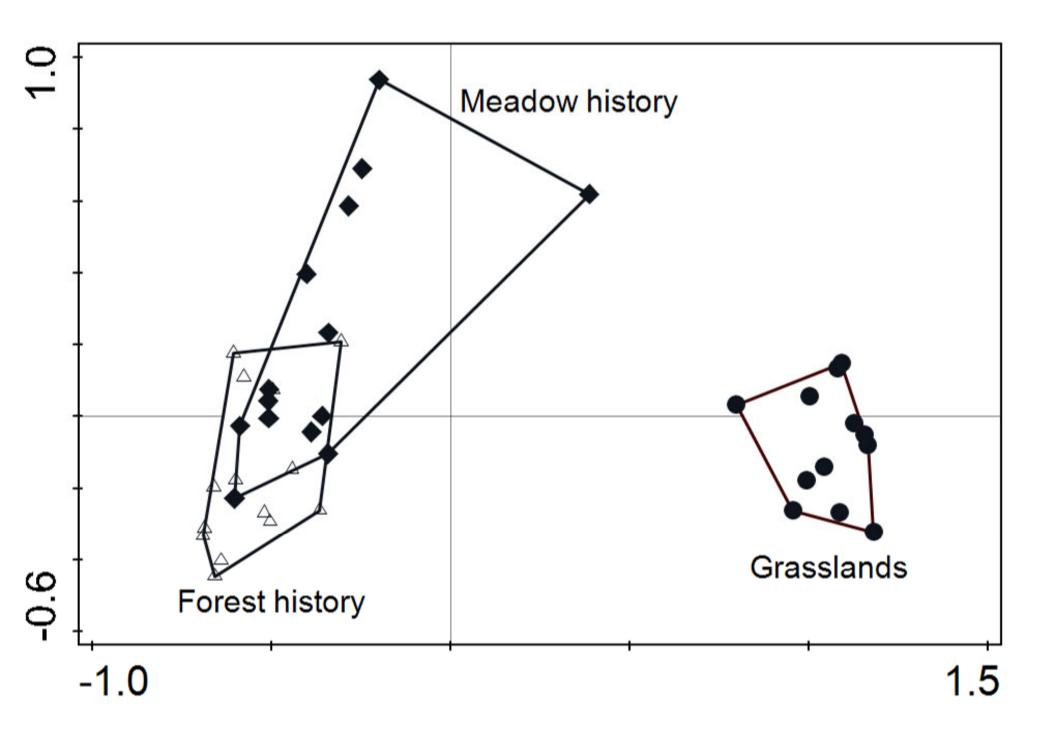


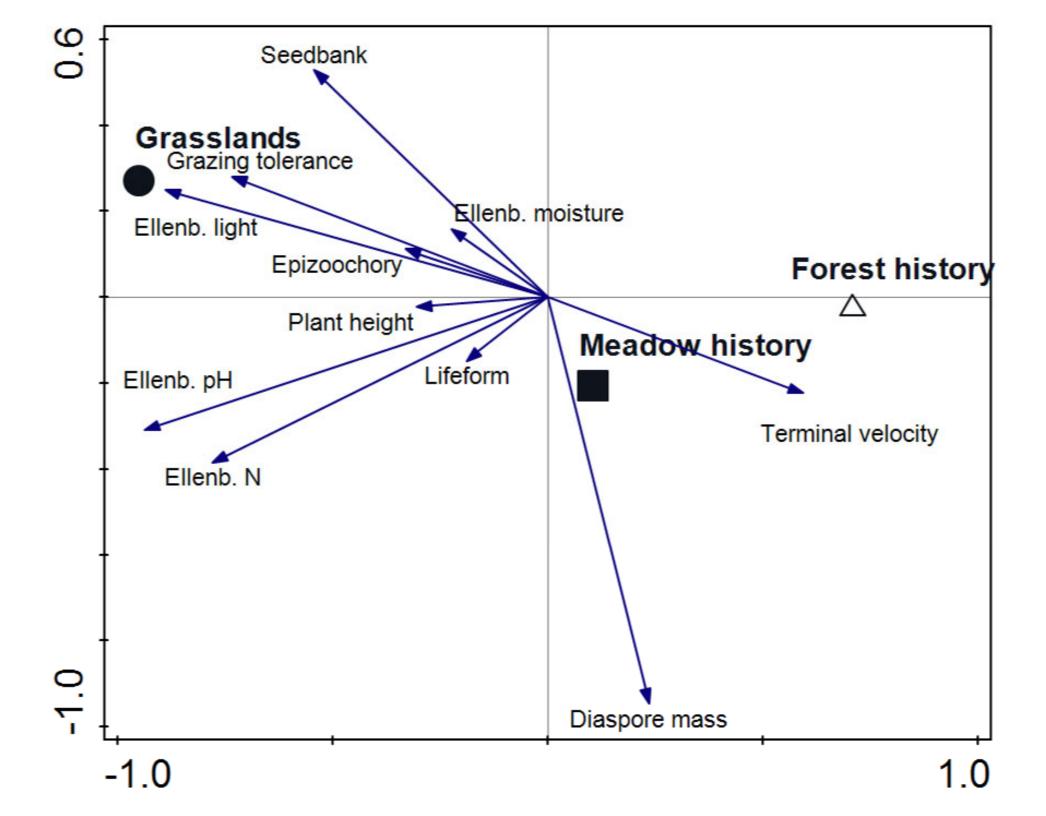
Equisetum pratense	
Geranium robertianum	
Rhododendron tomentosum	i
Vaccinium uliginosum	
Tussilago farfara	
Galeopsis tetrahit	
Stachys sylvatica	•
Cirsium arvense	
Poa nemoralis	
Phragmites australis	
Urtica dioica	_
Geum rivale	
Ranunculus repens	
Epilobium adenocaulon	
*Alchemilla vulgaris	•
Glechoma hederacea	
Senecio sylvaticus	-
Anthriscus sylvestris	_
*Pilosella lactucella	
Juncus effusus	
Aegopodium podagraria	
*Leucanthemum vulgare	-
Cirsium palustre	-
Phleum pratense	
Rumex acetosa	
*Scorzonera humilis	
*Ranunculus acris	
*Trollius europaeus	
Deschampsia cespitosa	
Astragalus glycyphyllos	
Juncuscon glomeratus	
_	
Prunella vulgaris Pyrola rotundifolia	
-	
Taraxacum sect.Ruderalia	
Daphne mezereum	
Anemone nemorosa	
*Filipendula ulmaria	
Gnaphalium sylvaticum	
Paris quadrifolia	
Equisetum arvense	
*Trifolium pratense	
Rubus saxatilis	-
Vicia cracca	
Lathyrus pratensis	_
Chamerion angustifolium	
*Pilosella officinarum	
*Veronica chamaedrys	
*Campanula rotundifolia	
*Primula veris	
Lactuca muralis	
Dryopteris filix-mas	
Clinopodium vulgare	_
Empetrum nigrum	
	-6 -4 -2 0 2

Anemone hepatica *Ajuga pyramidalis Trifolium medium	•
*Pimpinella saxifraga *Hypericum maculatum	
Agrostis capillaris	-
Hylotelephium telephium	
Equisetum sylvaticum Dactyils glomerata	
Stellaria media	
Achillea millefolium	
Solidago virgauera	-8-
Convallaria majalis Vicia sylvatica	
Hieracium sect.Hieracium	=
Knautia arvensis	
Geranium sylvaticum *Campanula persicifolia	
Rubus idaeus	-
Phegopteris connectilis	
Potentilla erecta	
Vicia sepium Fragaria vesca	
Viola riviniana	
Chrysosplenium alternifolium	
*Succisa pratensis	
*Lathyrus linifolius *Veronica officinalis	
Lycopodium annotinum	
Lysimachia vulgaris	
Calamagrostis arundinacea Melampyrum pratense/sylvaticum	
Polypodium vulgare	-
Maianthemum bifolium	=
Orthilia secunda Oxalis acetosella	
Potentilla palustris	
Luzula pilosa	
Trientalis europaea Mentha arvensis	
*Lotus corniculatus	
*Festuca ovina	-
*Anthoxanthum odoratum	
Calluna vulgaris Vaccinium vitis-idaea	•
Pteridium aquilinum	
Deschampsia flexuosa	
Vaccinium myrtillus Gymnocarpium dryopteris	
*Rumex acetosella	
Scutellaria galericulata	
Monotropa hypopitys	
Linaea borealis Goodyera repens	
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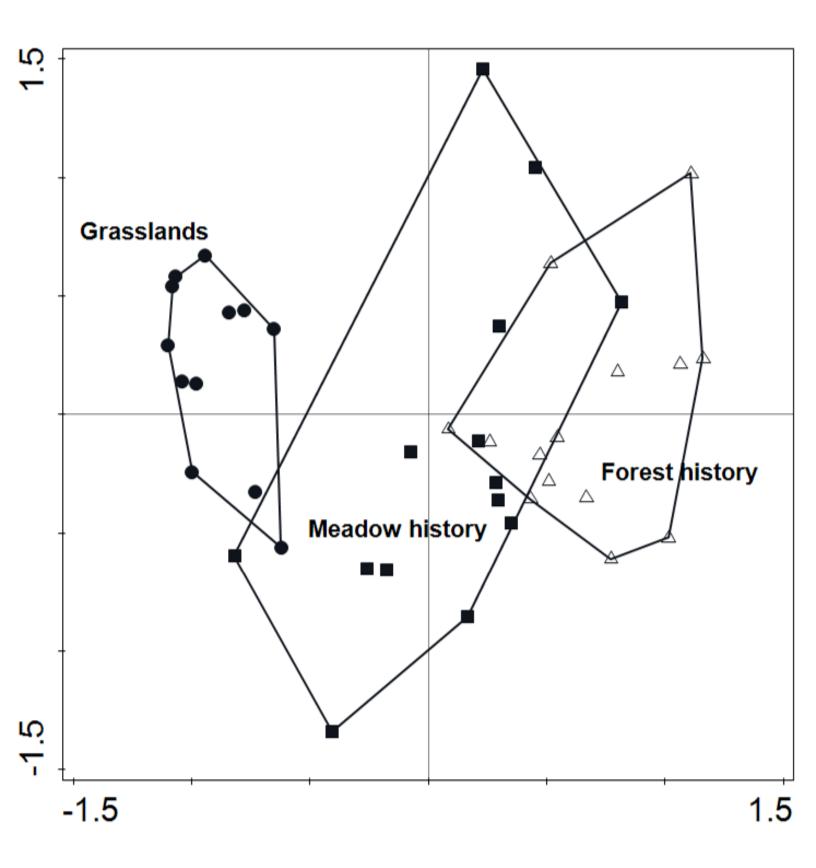


Table 1. Mean values and $CI_{95\%}$ for habitat attributes for the three different types of study sites: forest with a history as forest, forest with a history as meadow and grassland.

	History as forest		History as meadow		Grassland	
	Mean	Cl _{95%}	Mean	Cl _{95%}	Mean	Cl _{95%}
Size (ha)	4.26	3.21 - 5.31	3.98	3.26 - 4.70	4.42	3.57 - 5.27
Area with history as meadow (ha)			2.09	1.54 - 2.64		
Exposed soil (%)	6.75	3.30 - 10.21	10.64	6.71 – 14.57	2.03	0.33 – 3.73
Residue (%)	5.64	3.76 – 7.53	9.32	5.07 – 13.58	0.81	0.15 – 1.48
Bare rock (%)	1.34	0.94 – 1.74	1.39	0.82 – 1.95	6.46	4.24 - 8.68
Number of stumps (>10 cm)	0.55	0.40 - 0.71	0.92	0.57 - 1.26	0.13	0.06 - 0.19
Basal area coniferous (m²/ha)	22.20	18.88 – 25.51	19.23	15.29 – 23.16	0.10	0.03 - 0.24
Basal area deciduous (m²/ha)	0.76	0.29 - 1.23	1.76	0.29 – 3.22	2.56	1.43 - 3.68
Basal area total (m²/ha)	22.96	19.76 – 26.15	20.98	17.61 – 24.36	2.66	1.55 – 3.77

Traits	Variable type	Values	Proportion availble (%)	Source	
Lifeform	Categorical	1 - 4	99	www.lundsbotaniska.se	
Plant height	Continuous	0.053 - 2.50	97	Kleyer et al. 2008	square root transformed
Grazing/Moving tolerance	Categorical	1-3	92	www.lundsbotaniska.se	
Ellenberg light	Categorical	1-9	93	www.lundsbotaniska.se	
Ellenberg moisture	Categorical	1-10	85	www.lundsbotaniska.se	
Ellenberg pH	Categorical	1-9	65	www.lundsbotaniska.se	
Ellenberg nitrogen	Categorical	1-8	88	www.lundsbotaniska.se	
Diaspore mass	Continuous	0.002 - 43.9	90	Hintze et al. 2013, Kleyer et al. 2008	square root transformed
Terminal velocity	Continuous	0.075 - 2.45	83	Hintze et al. 2013, Kleyer et al. 2008	square root transformed
Epizoochory	Rank	0-1	65	Hintze et al. 2013	
Seed Bank	Categorical	1-4	72	www.lundsbotaniska.se	

Table 1. List of traits, their data type, range of values, proportion of species for which trait values existed in the databases and source for the trait values.

Table 3. List of the SD of sitewise average of traits, and weighted traits, in forest with a meadow history and grasslands; continuous conifer forests are included for reference. P-values for the comparison of SD in grasslands and forest with meadow history

		SD	-Cl _{95%}	+Cl _{95%}	Р
Lifeform					
SD	Grassland Meadow	0.669	0.371	0.966	ns
	history	0.632	0.374	0.890	
	Forest history	0.590	0.349	0.831	
Weighted SD	Grassland Meadow	0.440	0.244	0.636	ns +
	history	0.601	0.356	0.846	
	Forest history	0.621	0.368	0.840	
sqrt(plant height)	T OF EST HISTORY	0.021	0.508	0.875	
SD	Grassland	0.200	0.111	0.289	ns
50	Meadow	0.200	0.111	0.205	115
	history	0.211	0.125	0.298	
	Forest history	0.211	0.128	0.304	
Weighted SD	Grassland	0.175	0.097	0.253	ns
Weighted 5D	Meadow	0.175	0.057	0.233	115
	history	0.208	0.123	0.293	
	Forest history	0.208	0.123	0.293	
Grazing tolerance	rorest history	0.200	0.125	0.255	
SD	Grassland	0.748	0.415	1.081	ns
50	Meadow	0.740	0.415	1.001	115
	history	0.657	0.389	0.926	
	Forest history	0.631	0.373	0.888	
Weighted SD	Grassland	0.625	0.347	0.903	ns
Weighted 5D	Meadow	0.025	0.517	0.505	115
	history	0.565	0.334	0.795	
	Forest history	0.523	0.310	0.736	
Ellen light	,,				
SD	Grassland	1.473	0.817	2.128	ns
00	Meadow	1170	0.017	2.1220	
	history	1.758	1.040	2.475	
	Forest history	1.730	1.024	2.436	
Weighted SD	, Grassland	1.058	0.587	1.529	NS +
5	Meadow				
	history	1.732	1.025	2.439	
	Forest history	1.718	1.017	2.420	
Ellen moisture					
SD	Grassland	1.283	0.712	1.854	ns
	Meadow				
	history	1.236	0.732	1.741	
	Forest history	0.912	0.540	1.285	
Weighted SD	Grassland	1.026	0.570	1.483	ns
J	Meadow	-	-	-	
	history	0.960	0.568	1.351	
	Forest history	0.611	0.362	0.860	
Ellen pH	,				
SD	Grassland	2.097	1.164	3.030	ns
					-

	Meadow				
	history	2.040	1.207	2.873	
	, Forest history	1.919	1.136	2.703	
Weighted SD	, Grassland	1.860	1.033	2.688	ns
0	Meadow				-
	history	1.623	0.961	2.286	
	Forest history	1.301	0.770	1.832	
Ellen N					
SD	Grassland	2.080	1.154	3.006	ns
	Meadow				
	history	1.958	1.159	2.758	
	Forest history	1.835	1.086	2.584	
Weighted SD	, Grassland	1.877	1.042	2.712	ns
	Meadow				
	history	1.598	0.946	2.251	
	Forest history	1.445	0.855	2.034	
Diaspore					
SD	Grassland	0.972	0.539	1.405	ns
	Meadow				
	history	1.116	0.660	1.571	
	Forest history	1.224	0.724	1.723	
Weighted SD	, Grassland	0.935	0.519	1.351	ns
0	Meadow				-
	history	0.975	0.577	1.372	
	, Forest history	0.924	0.547	1.301	
sqrt(terminal					
velocity)					
SD	Grassland	0.413	0.229	0.597	ns
	Meadow				
	history	0.463	0.274	0.652	
	Forest history	0.426	0.252	0.600	
Weighted SD	Grassland	0.373	0.207	0.539	ns
-	Meadow				
	history	0.435	0.257	0.613	
	Forest history	0.354	0.209	0.498	
epizoo					
SD	Grassland	0.295	0.163	0.426	ns
	Meadow				
	history	0.318	0.188	0.448	
	Forest history	0.330	0.195	0.464	
Weighted SD	Grassland	0.290	0.161	0.419	ns
	Meadow				
	history	0.301	0.178	0.424	
	Forest history	0 277	0.164	0.390	
Seedbank	T OTEST HISTORY	0.277	0.104	0.550	
Jeeubank	Torest history	0.277	0.104	0.550	
SD	Grassland	1.052	0.164	1.520	ns
					ns
	Grassland				ns
	Grassland Meadow	1.052	0.584	1.520	ns
	Grassland Meadow history	1.052 1.069	0.584 0.632	1.520 1.505	ns ns
SD	Grassland Meadow history Forest history	1.052 1.069 1.073	0.584 0.632 0.635	1.520 1.505 1.511	
SD	Grassland Meadow history Forest history Grassland	1.052 1.069 1.073	0.584 0.632 0.635	1.520 1.505 1.511	
SD	Grassland Meadow history Forest history Grassland Meadow	1.052 1.069 1.073 0.941	0.584 0.632 0.635 0.522	1.520 1.505 1.511 1.359	

Appendix 1: Total plant species frequencies in three different study sites; grasslands, forests with a history as forest and forests with a history as meadow. Species classified as indicators for grasslands are marked with an "x" in the column to the right.

	Caralas	-		
Species	Grassland (max 1200)	History as forest (max 1400)	History as meadow (max 1400)	Indicator species
Achillea millefolium	788	2	4	
Achillea ptarmica	0	0	2	
Actaea spicata	0	0	1	
Aegopodium podagraria	63	2	23	
Agrimonia eupatoria	4	0	0	х
Agrostis capillaris	737	29	67	
Ajuga pyramidalis	53	9	24	x
Alchemilla vulgaris	620	0	8	х
Andromeda polifolia	0	0	1	
Anemone hepatica	34	62	170	
Anemone nemorosa	1	4	23	
Anthoxanthum odoratum	474	45	22	Х
Anthriscus sylvestris	295	7	95	
Aquilegia vulgaris	0	0	3	
Arctium minus	0	0	4	
Argentina anserina	6	1	0	х
Arnica montana	1	0	0	Х
Arrhenatherum elatius	1	0	4	
Artemisia vulgaris	5	1	0	
Astragalus glycyphyllos	0	0	3	
Bistorta vivipara	2	0	0	х
Briza media	101	0	0	х
Calamagrostis arundinacea	34	418	408	
Calluna vulgaris	21	160	81	
Campanula persicifolia	85	80	131	х
Campanula rotundifolia	376	3	10	х
Chamerion angustifolium	6	2	7	
Chelidonium majus	0	0	1	
Chrysosplenium alternifolium	5	3	4	
Cirsium acaule	1	0	0	х
Cirsium arvense	3	0	11	
Cirsium palustre	95	1	10	
Cirsium vulgare	13	0	2	
Clinopodium vulgare	12	3	9	
Convallaria majalis	20	27	48	
Crepis praemorsa	1	0	0	Х
Cynosurus cristatus	137	0	0	Х
Dactyils glomerata	212	17	35	
Dactylorhiza maculata	0	0	4	Х
Daphne mezereum	0	2	13	
Deschampsia cespitosa	142	16	116	
Deschampsia flexuosa	710	1048	759	

Dryopteris filix-mas	56	63	175	
Elytrigia repens	4	0	3	
Empetrum nigrum	0	1	3	
Epilobium adenocaulon	13	2	34	
Equisetum arvense	2	6	28	
Equisetum hyemale	0	12	0	
Equisetum palustre	1	0	3	
Equisetum pratense	20	1	90	
Equisetum silvaticum	33	65	130	
Euphrasia stricta var. stricta	14	0	0	Х
Festuca ovina	348	103	53	Х
Filipendula ulmaria	104	17	83	
Filipendula vulgaris	5	0	1	Х
Fragaria vesca	344	103	142	
Galeopsis tetrahit	7	0	14	
Gentianella campestris	2	0	0	х
Geranium robertianum	2	0	34	
Geranium sylvaticum	305	25	43	
Geum rivale	183	5	86	
Glechoma hederacea	40	0	7	
Gnaphalium sylvaticum	13	0	2	
Goodyera repens	0	28	0	
Gymnocarpium dryopteris	4	32	9	
Helianthemum nummularium	21	0	0	Х
Hieracium sect. Hieracium	310	90	151	
Hylotelephium telephium	2	1	2	
Hypericum maculatum	688	61	143	
Hypericum perforatum	2	0	1	Х
Inula salicina	1	0	0	
Juncus articulatus	5	0	0	х
Juncus bufonius	0	0	2	
Juncus conglomeratus	62	0	3	
Juncus effusus	57	3	36	
Knautia arvensis	379	19	33	
Lactuca muralis	16	110	289	
Lactuca serriola	0	0	1	
Lathyrus linifolius	525	260	303	Х
Lathyrus pratensis	349	6	21	
Leontodon autumnalis	124	0	0	Х
Leucanthemum vulgare	49	0	5	х
Linaea borealis	0	123	9	
Lolium perenne	6	0	0	
Lotus corniculatus	194	4	2	Х
Luzula pilosa	118	881	689	
Lycopodium annotinum	0	16	18	
Lycopodium clavatum	0	0	1	
Lycopus europaeus	0	0	1	
Lysimachia vulgaris	0	22	24	
Maianthemum bifolium	12	205	156	

Matricaria suaveolens	3	0	0	
Melampyrum pratense/sylvaticum	29	414	379	
Mentha arvensis	6	2	1	
Menyanthes trifoliata	0	0	1	
Monotropa hypopitys	0	5	0	
Myosotis scorpioides	5	0	1	х
Nardus stricta	40	0	0	х
Origanum vulgare	2	0	2	
Orthilia secunda	4	49	34	
Oxalis acetosella	25	585	452	
Paris quadrifolia	0	0	2	
Persicaria hydropiper	2	0	0	
Persicaria maculosa	1	0	0	
Peucedanum palustre	0	0	1	
Phegopteris connectilis	2	27	43	
Phleum pratense	120	1	10	
Phragmites australis	1	0	9	
Pilosella lactucella	4	0	6	Х
Pilosella officinarum	441	2	7	Х
Pimpinella saxifraga	306	4	10	Х
Plantago lanceolata	725	0	2	Х
Plantago major	129	0	0	
Plantago media	7	0	0	Х
Poa nemoralis	0	0	11	
Polygala vulgaris	71	0	0	Х
Polygonatum odoratum	1	0	0	
Polygonum aviculare	20	0	0	Х
Polypodium vulgare	6	100	75	
Potentilla argentea	4	0	0	Х
Potentilla erecta	569	129	185	
Potentilla norvegica	1	0	0	
Potentilla palustris	11	5	3	
Primula veris	61	3	10	Х
Prunella vulgaris	159	0	3	
Pteridium aquilinum	290	365	178	
Pyrola rotundifolia	1	0	3	
Ranunculus acris	785	9	71	Х
Ranunculus flammula	1	0	1	
Ranunculus repens	101	8	131	
Rhinanthus minor	33	0	0	Х
Rhododendron tomentosum	0	0	27	
Rubus chamaemorus	0	0	5	
Rubus idaeus	112	152	238	
Rubus saxatilis	37	33	138	
Rumex acetosa	578	2	18	v
Rumex acetosella	28	2	0	X
Satureja acinos	0	0	1	X
Saxifraga granulata	12	0	0	X
Scorzonera humilis	38	2	18	Х

Scrophularia nodosa	0	0	3	
Scutellaria galericulata	0	2	0	
Senecio sylvaticus	1	0	7	
Senicio viscosus	0	0	1	
Solanum dulcamara	0	0	1	
Solidago canadensis	0	0	1	
Solidago virgauera	8	44	82	
Stachys sylvatica	0	0	13	
Stellaria media	251	19	39	
Succisa pratensis	44	31	41	х
Taraxacum sect. Ruderalia	370	7	47	
Trientalis europaea	6	201	110	
Trifolium medium	9	3	8	
Trifolium pratense	761	8	36	х
Trifolium repens	504	0	1	х
Trollius europaeus	22	1	8	х
Tussilago farfara	5	0	24	
Urtica dioica	34	4	71	
Vaccinium myrtillus	206	1027	645	
Vaccinium oliginosum	1	0	25	
Vaccinium oxycoccos	0	0	7	
Vaccinium vitis-idaea	120	628	376	
Veronica chamaedrys	686	74	227	х
Veronica officinalis	215	143	164	х
Vicia cracca	429	7	29	
Vicia sepium	145	31	44	
Vicia sylvatica	0	5	9	
Viola riviana	503	303	391	
Total frequency	18569	8643	9419	
Total species richness	133	91	134	
Total richness indicator species	46	19	27	48
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