

Plant and soil responses to defoliation: a comparative study of grass species with contrasting life history strategies

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Abstract The overall aim of this study was to test for inter-species variation in plant and soil responses to defoliation among a broad range of temperate grass species and life-history strategies. We used a microcosm experiment where a range of grass species differing in life history traits were subjected to different intensities of defoliation, and a range of aboveground and belowground plant and soil responses were measured. All plant attributes, including accumulated shoot biomass, root biomass and root length, showed a strong negative response to defoliation, although plant species exhibited subtle differences in the way that they responded to increased severity of defoliation. Defoliation also exerted a strong influence on soil properties, decreasing soil microbial carbon (C) and the soil microbial C:nitrogen (N) ratio, and increasing inorganic N availability and potential N mineralisation across all species. Despite the wide range in life history strategies, plant species did not differ in their influence on most of the soil variables, except for the rate of nitrate mineralisation, which was lowest under plant species that displayed the least relative detrimental responses to defoliation. Collectively, our results suggest that plant

and soil responses to defoliation are reasonably consistent across a broad range of grass species, with only subtle inter-specific differences among species.

Keywords Defoliation · Microbial biomass · Grassland · Nitrogen cycling · Herbivory · Life history strategies

Introduction

The last few decades have witnessed an increasing attention being devoted to understanding the role that grazers play in regulating soil biogeochemical processes (Bardgett and Wardle 2010). This research has shown how grazing can modify a number of ecosystem properties, all of which ultimately impact on rates of soil nutrient cycling with feedback consequences for primary production. For example, grazing-induced increases in soil compaction (Cumming and Cumming 2003), alterations of soil water regimes (Medina-Roldán et al. 2007), plant species composition (Ritchie et al. 1998; Wardle et al. 2001), and aboveground and belowground primary productivity (Milchunas and Lauenroth 1993; Burke et al. 1998) are regularly cited as drivers of soil nutrient dynamics. Also, grazing animals can affect the spatial distribution of nutrients within ecosystems as a result of their movements and patchy return of nutrients in excreta, which in turn influences vegetation patterns (de Mazancourt et al. 1998; Bardgett and Wardle 2010).

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Over time scales from days to months, plant responses to biomass removal, or defoliation, act as a key mechanism by which grazing affects soil nutrient dynamics. Defoliation has been shown to alter plant allocation patterns, in terms of both root biomass (Mikola et al. 2001a) and root exudation (Paterson and Sim 1999, 2000). These changes in plant allocation can, in turn, have cascading effects on belowground food-webs, modifying microbial abundance and activity (Guitian and Bardgett 2000; Mikola et al. 2001a; Hamilton and Frank 2001) and the abundance of microbial predators (Mikola et al. 2001a; Hokka et al. 2004). All together, the effects on belowground food-webs mediated by defoliation have been linked to increases in soil nutrient mineralisation and availability (Hamilton and Frank 2001; Mikola et al. 2001a), and this is thought to contribute to the compensatory response of plant growth to herbivory (Hamilton and Frank 2001; Mikola et al. 2009).

Past studies suggest that plant and soil responses to defoliation vary across plant species and with the frequency of defoliation (Chapin and Slack 1979; Briske 1996; Guitian and Bardgett 2000; Klironomos et al. 2004; Ilmarinen et al. 2005; Gastal et al. 2010). Such differential responses of plant species to defoliation have been explained on the basis of plant functional traits which influence soil-resource dynamics, such as root exudates entering into the decomposers food-web (Holland et al. 1996; Mawdsley and Bardgett 1997; Mikola and Kytöviita 2002), and traits directly linked to the uptake of nutrients and competition for these with soil microbes such as root biomass and root length density (Chapin and Slack 1979; Oosterheld 1992). Most studies on this topic, however, have focused on a handful of plant species, and none, as far as we are aware, have comprehensively and simultaneously tested the responses of a broad range of plant species to defoliation. As a result, little is known about the differential response of grassland plant species and their associated soil microbial communities to defoliation, and this hampers our ability to draw generalizations about plant and soil responses to grazing in grassland ecosystems. Here, we redress this lack of knowledge by testing for inter-species variation in plant and soil responses to defoliation among a broad range of temperate grass species and life-history strategies. Specifically, we test the hypotheses that soil biological properties related to N cycling would be correlated with

the particular response of a plant species to defoliation (i.e. soil responses could be predicted from the defoliation response of the plants). This was done by measuring aboveground and belowground plant responses of a range of common UK grass species to different intensities of defoliation, and associated changes in soil microbial biomass and rates of N cycling, in a glasshouse microcosm experiment.

Material and methods

Experimental set-up

A range of seven common British grass species representing a spectrum of life histories including response to grazing and defoliation (see Table 1 for a summary of RGR_{max}, RGR_{mean}, C-S-R strategy, Ellenberg indicator values, and grazing and defoliation response), were grown in a glasshouse experiment during the spring of 2007 at Lancaster University, U.K. Briefly, seeds (Ermorsgate Seeds, Norfolk, UK) of the grass species *Nardus stricta* L., *Anthoxanthum odoratum* L., *Festuca rubra* L., *Poa pratensis* L., *Agrostis capillaris* L., *Lolium perenne* L., and *Holcus lanatus* L. (nomenclature follows Clapham et al. 1987), were germinated on a 1:1 (w/w) mixture of commercial acidic sand and Levingtons M3 growing media. The slow growing species *N. stricta* was sown 6 weeks before the rest of the grasses in order to reduce differences in plant size which are inherently dependent on the wide range in RGR_{max} employed in this study (Grime and Hunt 1975) (Table 1). The selected species represent a range of grasses with different plant traits which occur on grasslands subjected to different management intensities from infertile upland grasslands (*N. stricta*), *Agrostis-Festuca* and semi-improved grasslands (*A. odoratum*, *F. rubra*, *P. pratensis*, *A. capillaris*, *H. lanatus*) to highly productive pastures (*L. perenne*) (Rodwell 1992; Grayston et al. 2001).

Established seedlings were transplanted into pots (11 × 11 cm, height and diameter) following a similar design as that used by Guitian and Bardgett (2000). Pots' substratum was composed of a nutrient-poor soil (N%=0.6) coming from and acidic (pH in H₂O=4.5) soil taken from a semi-natural *Festuca ovina-Agrostis capillaris* grassland (National Vegetation Classification: U4b, Rodwell 1992) located in Littledeale,

Table 1 Some ecological traits of the species selected for the experimental study. Rmax=maximum relative growth rate (week⁻¹), Rmean=mean relative growth rate (week⁻¹)

Species	Rmax	Rmean	Ref	CSR-strategy	Ref	Ellenberg	Ref	Grazing response	Ref
<i>N. stricta</i>	0.71	0.71	1	S-SC	2	–		Increaser under sheep grazing	6
<i>A. odoratum</i>	0.94	0.94	1	SR-CSR	2	3	3	Increaser under rabbit grazing	5
<i>F. rubra</i>	1.18	1.18	1	C-S-R (but variable populations in C,S-C or S)	2	3	3	Decreaser under rabbit grazing	5
<i>P. pratensis</i>	1.26	1.26	1	C-S-R	2	6	4	Grazing tolerant	7
<i>L. perenne</i>	1.3	1.3	1	CR-CSR	2	7	3	–	
<i>A. capillaris</i>	1.36	1.36	1	C-S-R	2	4	4	Increaser under rabbit grazing	5
<i>H. lanatus</i>	2.01	1.56	1	C-S-R	2	5	4	Decreaser under rabbit grazing	5

References.

1. Grime and Hunt (1975)
2. Grime(2007)
3. Elberse and Berendse (1993)
4. Hill and Carey (1997)
5. Crawley (1990)
6. Welch (1986)
7. Hamilton and Frank (2001)

Lancashire, UK (Bardgett et al. 2003). Two weeks later, each grass species planted in the experimental pots was randomly assigned to 1 out of 3 defoliation treatments simulating different intensities of grazing, namely: 1) control plants without defoliation (UN); 2) light defoliated plants, clipped each 2 weeks (LD); and 3) heavy defoliated plants clipped at weekly intervals (HD). The defoliation treatment was imposed over an 8-week period and consisted of the removal of above-ground tissue at 4 cm above the soil surface. Treatments were applied in a randomised block design with 4 replicates per plant species and defoliation treatment combination, yielding a total of 84 pots (7 species X 3 defoliation X 4 blocks). Plants were watered on every other day and remained in a glasshouse in Lancaster University with the following settings: 12 h light-night periods providing a mean irradiance of 430 Wm⁻², mean day temperature of 22°C, and a mean night temperature of 18°C.

Measurement of aboveground variables

Before applying the defoliation treatments, initial shoot biomass was calculated in order to use this measure as a covariate for statistical analyses, and hence rule out size-specific effects of different plant species on response variables. This calculation was

based on species-specific allometric relationships between leaf length and shoot biomass of a subset of grass seedlings (*N. stricta*, $y = -6.79x^{1.2}$, $R^2 = 0.87$, $P < 0.001$, $n = 17$; *A. odoratum*, $y = -6.3x^{1.3}$, $R^2 = 0.97$, $P < 0.001$, $n = 6$; *F. ovina*, $y = -4.64x^{0.9}$, $R^2 = 0.96$, $P < 0.001$, $n = 18$; *H. lanatus*, $y = -5.37x^{1.1}$, $R^2 = 0.78$, $P < 0.001$, $n = 20$; *P. pratensis*, $y = -3.53x^{0.9}$, $R^2 = 0.95$, $P < 0.001$, $n = 17$; *A. capillaris*, $y = -7.45x^{1.2}$, $R^2 = 0.96$, $P < 0.001$, $n = 18$; and *L. perenne*, $y = -6.73x^{1.2}$, $R^2 = 0.93$, $P < 0.001$, $n = 17$). We, then, estimated leaf length in the experimental pots (digital images analysed in JMicroVision for Linux; Roduit 2007) and initial shoot biomass was calculated based on the estimated leaf lengths using the aforementioned allometric parameters. At the end of the experiment, all aboveground plant biomass was harvested from each pot and combined with that collected from previous clippings to determine total shoot biomass (after 70°C drying for 48 h) produced over the experimental period.

Measurement of belowground variables

Plant root systems were manually removed from each pot to estimate root-related variables. Total root length was estimated by species-specific allometric relationships between root length and root mass from root

sub-samples. First, we scanned sub-samples of the fresh root systems in a conventional scanner (CanoScan 4200 F, Canon Inc. Tokyo Japan). Roots were submerged into water in an acrylic tray avoiding root overlapping as much as possible and scanned in a 300 dpi resolution. These digital images were analysed using an image analysis software (ImageJ, Rasband 2007) by the method described in Kimura et al. (1999), and this estimation of root length was used to model total root length using total root mass as the predictor. This approach produced highly significant parameters to estimate total root length for all species (data not included). Root dry biomass (including grass crowns) was determined by oven-drying the roots as with aboveground biomass.

Soil collected after root recovery was passed through a 2 mm mesh sieve and stored at 4°C before the analytical assays were carried out. Microbial biomass carbon (C) and N were determined by extracting 5 g of soil (fresh basis) in 25 mL of 0.5 M K₂SO₄ of both 24 h-CHCl₃ fumigated and unfumigated soil samples (Vance et al. 1987). Soil-K₂SO₄ extracts were shaken for 30 min and filtered through a 1-grade Whatman paper. Microbial biomass C was calculated as the difference in total C in fumigated and unfumigated soil extracts, determined with a Shimadzu 5000A TOC analyser (Shimadzu Inc., Japan) and using an extraction efficiency factor (ke_C) of 0.45 (Sparling et al. 1990). Microbial biomass N was measured as the difference in total N (N-NH₄⁺ and N-NO₃⁻) produced by the persulfate digestion method (Cabrera and Beare 1993) between fumigated and unfumigated soil extracts, measured by automated continuous flow colorimetry in a Bran and Luebbe AutoAnalyser 3. The extraction efficiency factor (ke_N) used for N in microbial biomass was 0.54 (Brookes et al. 1985).

Potential NH₄⁺ mineralisation rate was determined by measuring the N-NH₄⁺ difference between KCl extracts of incubated and non-incubated soil samples (Campbell et al. 1993; Harrison and Bardgett 2010). Non-incubated samples were extracted without previous incubation, whereas incubated samples were held for a period of 14 days in sealed sterile bottles at 25°C. Extractions were made with 10 g of fresh soil extracted in 50 ml 1 M KCl, shaken for 60 min and filtered through a 1-grade Whatman paper. Ammonium mineralisation rate was calculated as $(N-NH_4^+ t_{14} - N-NH_4^+ t_0)/(14 \text{ days})$ (mg N-NH₄⁺ kg dry soil⁻¹ day⁻¹).

Nitrate mineralisation rate was determined following the same procedure as for N-NH₄⁺, but for N-NO₃⁻. Both N-NH₄⁺ and N-NO₃⁻ in KCl extracts were measured by automated continuous flow colorimetry as described above.

Statistical analyses

All variables were tested for normality with the Shapiro-Wilks test and normalised if necessary (Zar 1998). Data were analysed using ANOVA with plant species and defoliation as fixed experimental factors, and blocks as a random factor by using general linear mixed models (Pinheiro et al. 2008). For plant-related variables (aboveground and belowground biomass, root length, and specific root length), ANCOVA was used to test the effect of plant species and defoliation including initial shoot biomass as a covariate term and models' estimates were corrected depending on the significance of these term (Engqvist 2005). Estimated shoot initial biomass (see aboveground variables) values were (average dry weight (mg)±standard deviation): *N. stricta* (22±15), *F. rubra* (60±12), *P. pratensis* (160±43), *H. lanatus* (310±39), *A. odoratum* (373±134), *A. capillaris* (16±3), and *L. perenne* (165±49); thus justifying its use in the ANCOVA models. All statistical tests were carried out with the R statistical package for Linux (R Development Core Team 2010).

Results

Aboveground responses

At final harvest, the extent that total accumulated shoot biomass was reduced by defoliation varied among the grass species, as shown by the significant defoliation by species interaction ($F_{12,57}=7$, $P<0.0001$, Fig. 1a). Relative reductions in total accumulated shoot biomass resulting from defoliation were more pronounced for *F. rubra*, *H. lanatus*, *A. odoratum* and *A. capillaris* when defoliation intensity was low (72, 56, 65 and 57% reductions in shoot biomass between lightly-clipped and undefoliated controls, respectively) than under more severe defoliation (6, 20, 7 and 38% reductions in shoot biomass between heavy and light defoliation, respectively). In contrast, decreases in shoot biomass of *P. pratensis* and *N. stricta* were of a similar magnitude

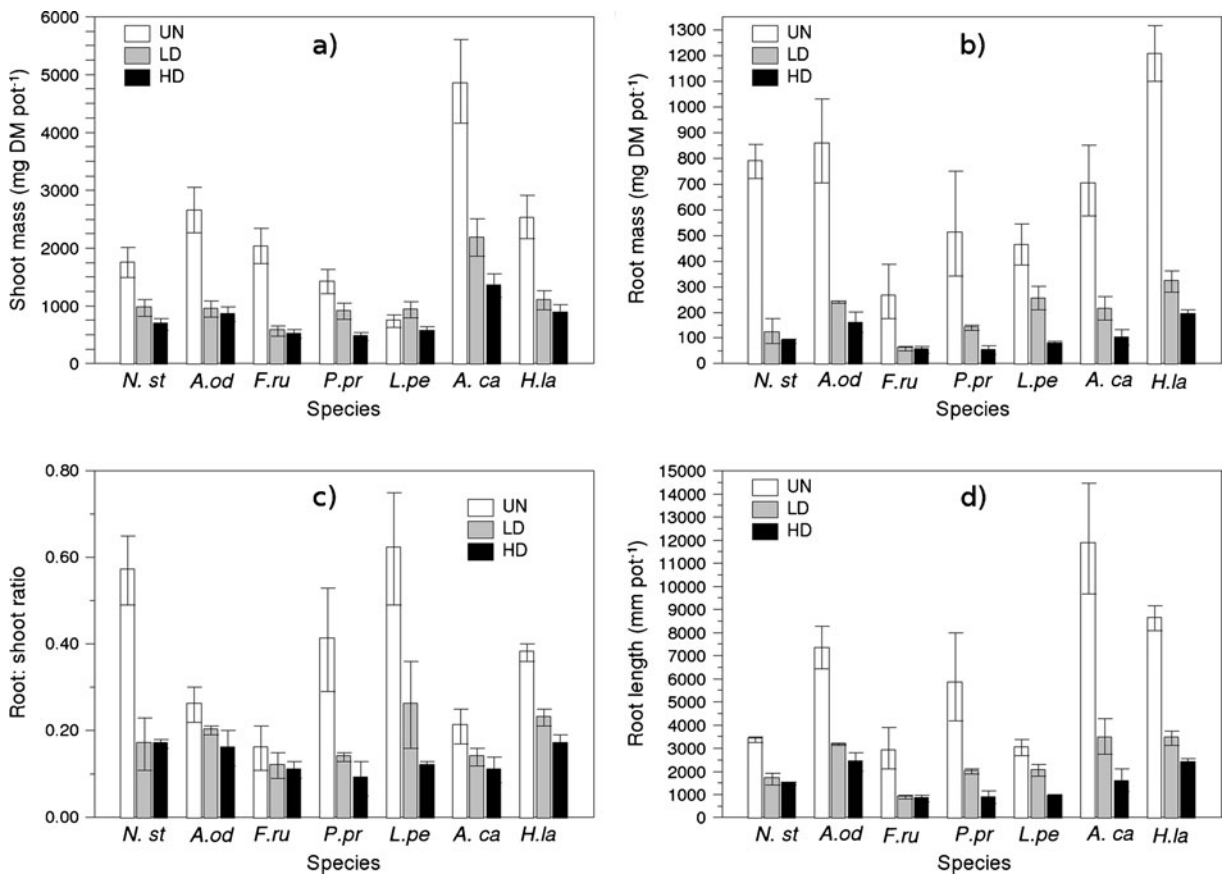


Fig. 1 Plant response to different intensities of defoliation across a broad range of common grass species, including: **a** Total accumulated shoot biomass; **b** belowground biomass; **c** root: shoot ratio; and **d** total root length (as estimated from allometric equations fitted to root mass and root length data estimated by image analysis). Labels are: UN=undefoliated controls ; LC=light defoliation (defoliation each two weeks); and HD=heavy defoliation (weekly). *N.st*=*Nardus stricta*, *A.*

od=*Anthoxanthum odoratum*, *F.ru*=*Festuca rubra*, *P.pr*=*Poa pratensis*, *L.pe*=*Lolium perenne* *A.ca*=*Agrostis capillaris*, *H.la*=*Holcus lanatus*. All values are back-transformed means±1 SE. In the case of aboveground biomass, values are adjusted means±1 adjusted SE as determined by ANCOVA with biomass in the beginning of experiment being the significant covariate

across the defoliation frequency treatments. Finally, shoot biomass of *L. perenne* increased under light defoliation but decreased moderately under heavy-defoliation frequency. Individual ANOVA's for each grass species (note that cross-species comparisons are not valid for these analyses) showed that species roughly grouped into those whose shoot biomass significantly decreased for both defoliation frequencies (i.e., *A. capillaris*, *H. lanatus* and *P. pratensis*), those which were sensitive to light but not further high intensify defoliation (i.e., no difference between lightly- and heavily-defoliated plants, namely *A. odoratum*, *F. rubra* and *N. stricta*), and *L. perenne* which, although responsive to defoliation, did not show a clear

separation among defoliation intensities because it yielded a very low shoot biomass in the undefoliated controls (Fig. 1a).

Belowground responses

Root biomass decreased strongly in response to defoliation across all plant species ($F_{2,55}=140.5$, $P<0.0001$, Fig. 1b). Unlike shoot biomass, the decrease in root biomass in response to defoliation was similar for all species (species X defoliation interaction effect, $F_{12,55}=1.5$, $P=0.17$), and all species showed a high sensitivity to light defoliation (an average 71% reduction between light defoliation and controls).

Post-hoc tests showed that there were no differences between the means for the light- and heavy-frequency defoliation for *A. odoratum*, *F. rubra*, and *N. stricta* (data not shown). Plant species significantly differed in root biomass ($F_{6,55}=18$, $P<0.0001$, values for the individual q statistics of the Tukey's test not included) in the order $H. lanatus > A. odoratum > A. capillaris$, $N. stricta = L. perenne > P. pratensis > F. rubra$.

We detected a significant defoliation by species interaction for the root to shoot mass ratio (R:S) ($F_{12,55}=3$, $P=0.004$) which provides additional evidence that plant species responded differently across the two defoliation frequencies (Fig. 1c). No clear relationship could be observed for the reductions in R:S in response to defoliation or in relation to species life history strategy, at least based on published values of RGR_{max} .

Total root length (TRL) decreased in response to defoliation (Fig. 1d), and this reduction varied among the plant species with defoliation frequency as shown by the species by defoliation interaction ($F_{12,52}=2$, $P<0.05$). *A. capillaris* and *P. pratensis* showed similar reductions in TRL across the two defoliation frequencies (*A. capillaris*=71% reduction between the undefoliated control and light defoliation, and 55% between light defoliation and the heavy defoliation treatment; *P. pratensis*=65 and 57% for the same treatments respectively). On the other hand, the reduction in TRL of *L. perenne* was higher when defoliation frequency increased (33% reduction between undefoliated controls and light defoliation and 55% reduction between light and heavy defoliation respectively). *A. odoratum*, *F. rubra*, *H. lanatus* and *N. stricta* were relatively insensitive to heavy defoliation, displaying only a low reduction in TRL from the low to the high intensity treatment. Individual ANOVA's for each grass species (data not shown) showed basically the same patterns as those already described, i.e., a group of species which responded consistently negative to increasing frequency of defoliation (i.e., *A. capillaris*, *H. lanatus*, *L. perenne* and *P. pratensis*) and another which was sensitive to low, but not further high intensity defoliation (as shown by the means differences between these treatments, *A. odoratum*, *F. rubra*, and *N. stricta*). Since we used root mass in our species-specific estimations of TRL, statistical analysis of specific root length would produce more or less the same qualitative results as those for root mass. However, results among species can portray some insight on the extent of the

costs species incurred in terms of root production (with lower values indicating higher construction costs). Specific root length followed the sequence $L. perenne (110\pm 10) < F. rubra (137\pm 11) < P. pratensis (141\pm 14) < N. stricta (143\pm 22) < H. lanatus (169\pm 14) < A. odoratum (172\pm 11) < A. capillaris (172\pm 12 \text{ mg}^{-1})$ respectively).

We used principal component analysis (PCA) on transformed response variables (aboveground and belowground biomass, root to shoot ratio and total root length) to summarize plant species responses to defoliation. The first PCA axis explained 78% of the variance in the data and was associated with defoliation frequency, with belowground plant attributes (i.e. root mass and total root length) having the highest loads on this first component (data not shown).

Soil responses

When data were integrated across all species, light and heavy frequency defoliation reduced microbial biomass C in soil ($F_{2,53}=7.5$ $P<0.01$, Fig. 2a). Microbial biomass C was reduced from 2040 ± 147 (mean \pm 1 SE) mg C kg⁻¹ in soil of control plants to 1480 ± 71 and 1465 ± 102 mg C kg⁻¹ in soil of plants under moderate and heavy defoliation frequency respectively and this reduction in microbial C was correlated to reductions in root biomass caused by defoliation ($r=0.36$, $t_{74}=3.5$, $P=0.001$). Post-hoc analyses showed, however, that microbial C means across defoliation frequencies were significantly different only for *H. lanatus* and *A. capillaris*, and for *A. odoratum* between the undefoliated and the clipped-treatments (Fig. 2a). In contrast, defoliation did not affect microbial biomass N ($F_{2,54}=1.5$, $P=0.26$, Fig. 2b); hence, across all plant species, microbial C:N ratio was reduced by 20.0 and 25.6% under moderate and heavy defoliation, respectively, when compared with the undefoliated control ($F_{2,53}=8.5$, $P<0.001$, Fig. 2c).

Soil N availability was also affected by defoliation treatments (Fig. 3a). Across all plant species, heavy defoliation increased soil NH_4^+ ($F_{2,54}=3$, $P=0.05$) and total inorganic N ($F_{2,54}=3$, $P=0.05$) concentrations by 35% and 29% respectively compared to the undefoliated control. Mean comparison tests showed significant differences in *N. stricta* and *A. capillaris* between undefoliated controls and the defoliated treatments but not in the rest of the species. Soil

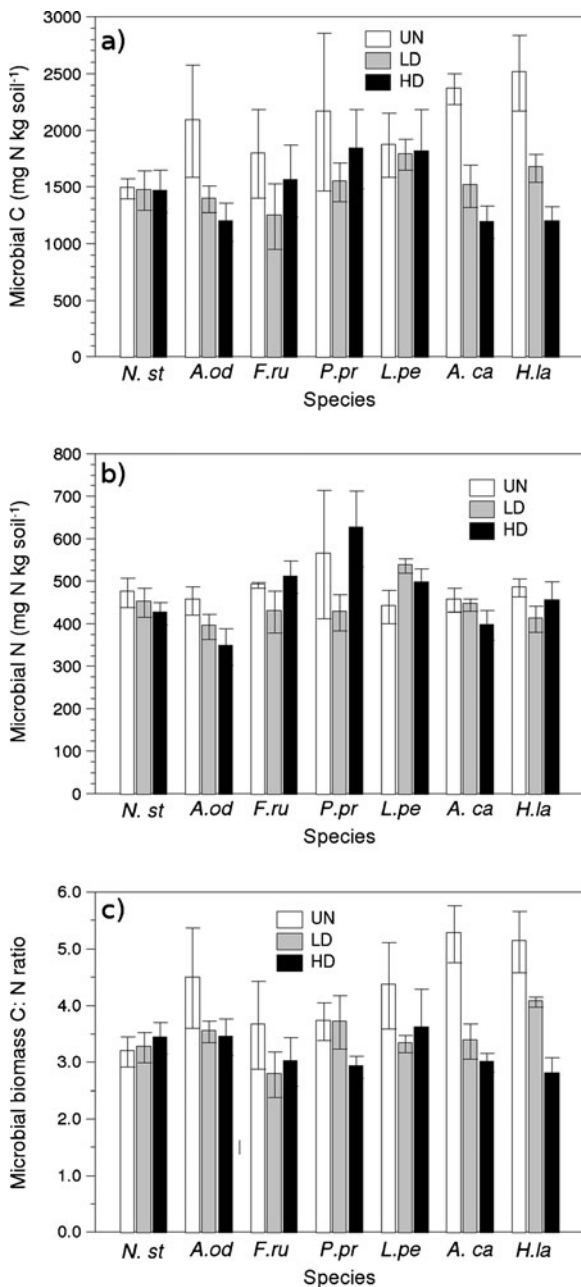


Fig. 2 Responses to different intensities of defoliation across a broad range of common grass species of soil microbial biomass carbon (a); soil microbial nitrogen (b); and the microbial C: N ratio (c). UN=undefoliated controls; LD=light-defoliation (defoliation each two weeks); HD=heavy defoliation (weekly). *N.st*=*Nardus stricta*, *A.od*=*Anthoxanthum odoratum*, *F.ru*=*Festuca rubra*, *P.pr*=*Poa pratensis*, *L.pe*=*Lolium perenne*, *A.ca*=*Agrostis capillaris*, *H.la*=*Holcus lanatus*. All values are means \pm 1 SE

NO_3^- concentrations were not significantly affected by defoliation ($F_{2,54}=3$, $P>0.05$, Fig. 3b), although potential net nitrification rate increased by almost 200% under heavy defoliation ($F_{2,54}=3.5$, $P<0.05$) relative to the undefoliated controls (Fig. 3c). The rate of net NO_3^- mineralisation also varied significantly among plant species ($F_{6,54}=3.5$, $P<0.01$, Fig. 3c), being greatest in soil planted with *N. stricta* (2.9 ± 1.9 mg NO_3^- kg soil⁻¹ day⁻¹) and lowest in soil of *A. capillaris* (0.21 ± 0.22 mg NO_3^- kg soil⁻¹ day⁻¹). The sequence of NO_3^- mineralisation after post-hoc tests was, in descending order, *N. stricta*, *L. perenne*, *F. rubra*, *P. pratensis*, *H. lanatus*, *A. odoratum*, and *A. capillaris*. Some plant species at the fast-growing end of the life history gradient had the lowest values of net NO_3^- mineralisation, but the overall trend was erratic (non significant species X defoliation interaction). Post-hoc tests showed that mean differences in NO_3^- mineralisation between heavy defoliation and the other treatments were significant for *A. odoratum*, *F. rubra*, *H. lanatus* and *P. pratensis*. Neither net potential NH_4^+ nor total N mineralisation rates (Fig. 3d) were affected by the experimental treatments and grass species identity did not, in general, influence soil variables, aside potential net NO_3^- mineralisation rate as already stated.

Discussion

In this study, we tested whether plant responses to defoliation vary across a range of grassland species representing a broad range in life-history strategies, and whether soil microbial and biogeochemical responses to defoliation were related to such inter-species variation. We detected a general detrimental effect of defoliation on shoot and root growth among all plant species tested, and subtle inter-specific differences in the response to defoliation frequency. In particular, *A. capillaris* and *H. lanatus* showed consistently higher absolute values in shoot and root biomass and root length, and relatively lower losses in root attributes across defoliation treatments (the relative change in biomass in Ferraro and Oesterheld 2002), but their performance was reduced when defoliation was intensified (from light to heavy defoliation frequencies). The same reductions in response to defoliation were observed in *P. pratensis* and *L. perenne*, but these later species displayed low

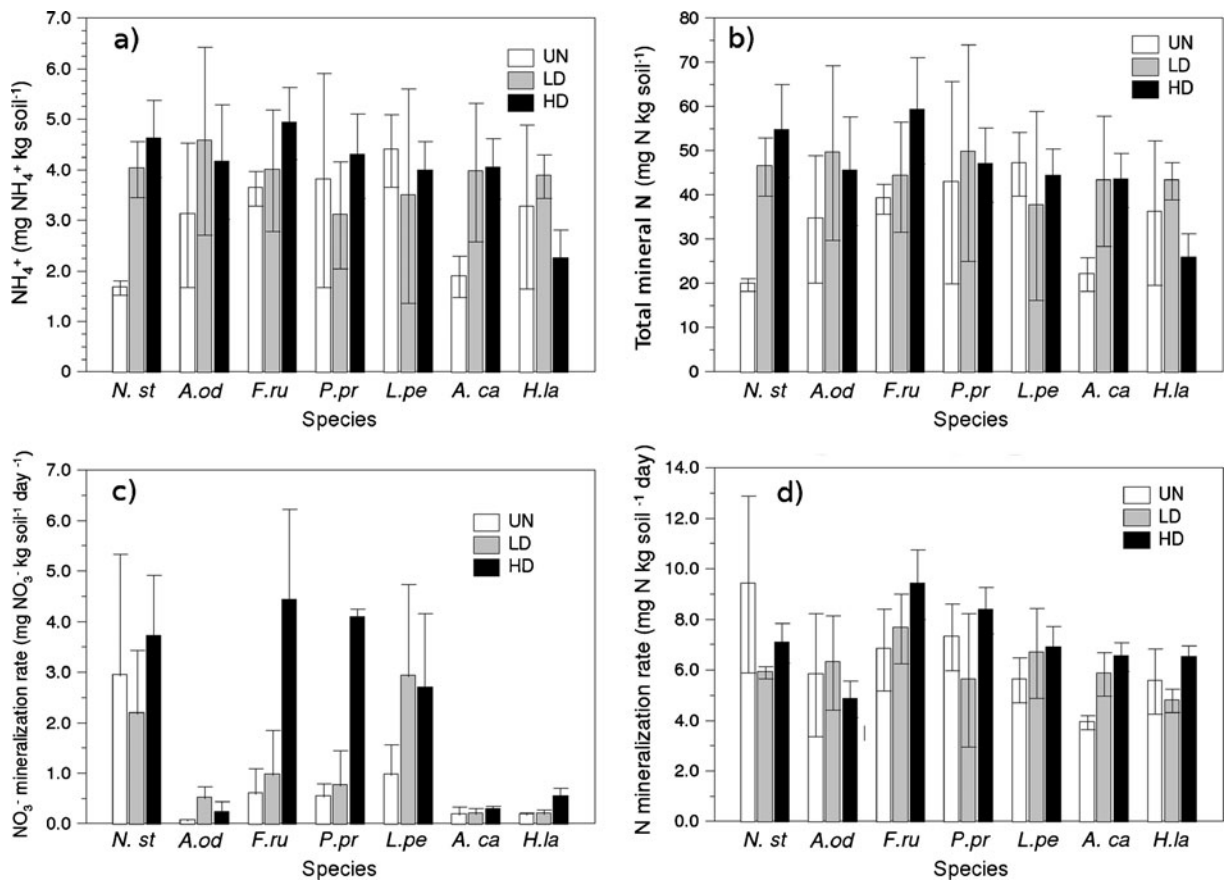


Fig. 3 Soil responses to different intensities of defoliation across a broad range of common grass species, including ammonium availability (a), mineral soil N (ammonium+nitrate) (b), net potential nitrate mineralisation rate (c) and the total N mineralisation rate (ammonium+nitrate) (d); UN=undefoliated

controls; LD=light defoliation (defoliation each two weeks); HD=heavy defoliation (weekly). *N.st*=*Nardus stricta*, *A.od*=*Anthoxanthum odoratum*, *F.ru*=*Festuca rubra*, *P.pr*=*Poa pratensis*, *L.pe*=*Lolium perenne*, *A.ca*=*Agrostis capillaris*, *H.la*=*Holcus lanatus*. All values are means ± 1 SE

absolute values in plant measures. Yet another group of species, *A. odoratum*, *F. rubra* and *N. stricta*, showed no additional decreases in performance between light and heavy defoliation in most of our plant measures, which we interpret here as a sign of resistance to defoliation. Abundance of *N. stricta* is often high in grazed grasslands (Welch 1986), which is attributed to its unpalatable shoot tissue (Massey et al. 2007), and Hartley and Amos (1999) reported that defoliation (a less severe regime than that used here) did not cause reductions in root length of *N. stricta* plants, adding evidence that this species is resistant to defoliation. There is evidence of *F. rubra* showing compensatory growth responses to grazing as well (Berg et al. 1997; Van der Graaf et al. 2005). Several other studies have likewise found declines in

root productivity in response to defoliation (Guitian and Bardgett 2000; Mikola et al. 2001a), although evidence is mixed. For example, fenced exclusion studies on Serengeti grasslands show that mammalian grazers do not necessarily inhibit root biomass and productivity (McNaughton et al. 1998), and in a global literature synthesis Milchunas and Lauenroth (1993) reported both enhancements and reductions in root biomass as a result of herbivore exclusion. Moreover, in a meta-analysis on graminoids, Ferraro and Oesterheld (2002) showed that the effects of defoliation were less acute on root than on shoot biomass. Nevertheless, our analysis showed that root mass and root length are important attributes in describing inter-specific differences in response to defoliation. Root length is a measure of plant foraging

scale (Kembel and Cahill 2005) and therefore it might be associated with the response of grasses to defoliation.

In general, the detrimental response of plant growth to defoliation was mirrored in the soil biological properties measured, although inter-specific differences were not detected. Across all plant species, defoliation was found to reduce microbial biomass and its C: N ratio, and to increase NH_4^+ availability and the rate of NO_3^- mineralisation. The negative response of microbial biomass C to defoliation that we observed contrasts sharply with results of a number of experimental studies, which have found that defoliation stimulates soil microbes (Mawdsley and Bardgett 1997; Bardgett et al. 1998; Mikola et al. 2001b). Increases in microbial biomass following defoliation have been attributed to the stimulation of root exudation (Holland 1995; Holland et al. 1996; Mawdsley and Bardgett 1997; Hamilton et al. 2008), and this stimulation of root exudates as a result of clipping has been reported for some of the plant species used in our experiment, namely *L. perenne* and *F. rubra* (Paterson and Sim 1999, 2000). Our findings indicate, however, that defoliation caused soil microbes to become limited by C, as evidenced by the decline in the microbial C:N ratio, which is indicative of increase in C relative to N limitation (Kaye and Hart 1997). This decline is also likely to be related to the reduction in root biomass across all species as a result of defoliation, a view supported by the positive correlation of root biomass with microbial biomass C. The lack of effect that increasing the intensity of defoliation had on microbial biomass C is likely explained by the relatively large effect that light defoliation had on plant performance relative to the undefoliated controls. The decline in microbial biomass C could also be partly due to increased predation by soil animals, given that previous studies have shown that defoliation enhances the abundances of microbial-feeding faunal groups in soil (Mikola et al. 2001a; b). However, in our study, it is most likely that reductions in root C allocation under defoliation, and hence C supply to soil, is the main cause of the consistent decline in microbial biomass across all defoliated plant species (Mikola et al. 2001a; Bazot et al. 2005; Hamilton et al. 2008; Sankaran and Augustine 2004).

Defoliation increased NH_4^+ and total inorganic N availability, as well as the potential rates of NO_3^-

mineralisation across all species tested. A number of studies have documented a stimulatory effect of defoliation (Hamilton and Frank 2001; Mikola et al. 2001a; Ayres et al. 2007) and ungulate grazing (Seagle et al. 1992; Hamilton et al. 2008) on soil N availability and mineralisation, and this response is thought to be a key mechanism contributing to compensatory growth in grazed grassland (Owen 1980; Ritchie et al. 1998; Hamilton and Frank 2001; Bardgett and Wardle 2003). Such enhanced soil N mineralisation has been attributed to a variety of mechanisms, including the return of N-rich plant litter and animal wastes to soil (Day and Detling 1990; de Mazancourt et al. 1998), and the stimulation of microbial activity and N mineralisation in the root zone due to enhanced root exudation in defoliated plants (Hamilton and Frank 2001; Mikola et al. 2001a, b). In our experiment, however, we propose that the defoliation-induced increase in soil N availability and mineralisation was due to the previously mentioned switch to C limitation of the microbial biomass, as indicated by the reduction in microbial C:N across all defoliated plants. Indeed, it is well established that under conditions when microbial growth is C limited, microbes use the C to support their energy needs and they excrete plant available ammonium (NH_4^+) as a waste product into soil i.e. N is mineralised by the microbial biomass (Kaye and Hart 1997). Microbial C limitation together with the strong negative impact of defoliation on plant size and the concomitant reduction in total plant N might be the cause for the increase in N availability. As previously mentioned, it is also possible that higher rates of microbial predation in soils of defoliated plants contributed to the stimulation of soil N availability via the microbial-loop (Clarholm 1985), although this was not measured in this study.

Despite the wide spectrum in ecological traits in the plant species we used, and the subtle differences in plant growth responses to defoliation, few inter-specific differences were observed in the response of soil properties. Only for the rate of NO_3^- mineralisation did we detect inter-specific differences in the responses to defoliation. Here, we found that the rate of NO_3^- mineralisation was significantly lower in soils planted with the grasses which showed the highest biomass values, namely *A. capillaris*, *A. odoratum* and *H. lanatus*, than in soils planted with

N. stricta. However, we did not detect a defoliation by species interaction, indicating that such inter-specific differences in NO_3^- mineralisation were independent of defoliation. This trend of lower rates of NO_3^- mineralisation in soils planted with those species which exhibited higher biomass values across defoliation treatments is difficult to interpret given that no concomitant changes in NH_4^+ availability were detected for the same set of species. The absence of inter-species differences in other soil properties across the species tested is in contrast to previous studies which show that plant species, and even genotypes, can have markedly different effects on soil biological properties, acting as major determinant of microbial communities in soil (Bardgett et al. 1999; Innes et al. 2004; Bezemer et al. 2006; Markham et al. 2009; Harrison and Bardgett 2010; Orwin et al. 2010). We do not know the reason for the absence of such inter-species variation in soil properties in our experiment. However, given that inter-specific differences in most soil biological properties were apparent in the undefoliated controls, but not in defoliated plants, it appears that defoliation has cancelled out any differences at the species level. Secondly, it has been shown that the effects of plant species on soil properties are dependent on soil type (Innes et al. 2004; Marschner et al. 2004; Bezemer et al. 2006). In this way, it seems that our soil might have restricted the expression of strong plant effects on soil biological properties, suggesting that other factors, such as low pH and nutrient availability, might have been primary determinants of these measures. Despite this, defoliation was found to consistently and strongly promote soil nutrient availability in soil across all species tested.

In conclusion, our results show that grassland plant species representing a broad range of life history strategies respond in a consistent way to defoliation. Across all species tested, we found that defoliation reduced plant growth, especially of root mass and length, but stimulated N availability in soil. We did not measure the consequences of this defoliation-induced stimulation of N availability, but we propose that it would, in the long term, positively feedback to the plant in terms of improved N acquisition and, potentially, improved growth. Surprisingly, we found only subtle differences in the response of different plant species to defoliation, and no inter-species variation in the response of soil properties to this treatment. This suggests that, in these soils, effects of

defoliation on plant and soil properties are remarkably consistent, at least under the experimental conditions used in this study. Overall, our findings reinforce the view that aboveground and belowground components of ecosystems are strongly interrelated and that understanding the effects of grazing on ecosystems nutrient cycling requires a combined aboveground and belowground approach.

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