Long-term belowground effects of grassland management: the key role of liming

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Long-term belowground effects of grassland management: the key role of liming

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Abstract

The functioning of human-managed grassland ecosystems strongly depends on how common management practices will affect grassland 'belowground compartment' including soil biogeochemistry and plant roots. Key questions remain about how animal grazing, liming (e.g. the addition of CaCO_3 to soils), and nutrient fertilization might affect, in the long-term, soil nutrient cycling and multiple root traits. Here we focus on a mesotrophic grassland located in Berkshire, UK, where contrasting levels of rabbit grazing, liming and different inorganic fertilizers have been applied since 1991. We ask how (1) soil nitrogen (N) availability and cycling, (2) total root mass, (3) root mass decomposition, and (4) arbuscular mycorrhizal fungal (AMF) root colonisation might respond to 22 years of very different management. We found that liming strongly affected total root mass, root decomposition, root AMF colonisation as well as soil N availability and cycling and that these effects were mainly driven by liming-induced increases in soil pH. Increases in soil pH were associated with significant (1) decreases in root mass, (2) increases in root mass decomposability and in the mineralization of N in decomposing root detritus, and (3) increases in AMF infection. Soil pH was also significantly related to greater N availability (i.e. soil NO_3 levels) and to lower $\delta^{15}\text{N}$ natural abundance, which suggests more efficient N uptake by plants in limed soils as we found in our study. The application of multiple nutrients (NPKMg) also reduced total root mass, while N-only fertilization was associated with greater AMF infection. Surprisingly the long-term impact of grazing was generally weak and not significant on most plant and soil parameters. Despite soil pH affecting most belowground variables, changes in soil pH were not associated with any change in soil C and N stocks. Because liming can improve nutrient cycling (and benefits soil pH and grass yields) without negatively affecting soil C sequestration, we suggest that regular liming applications may provide management solutions for increasing the long-term sustainability of permanent grassland.

Key Words: carbon sequestration, ecosystem services, grassland management, grazing, nitrogen cycling, nutrient fertilization, soil pH

Introduction

The functioning of human-managed grassland ecosystems greatly depends on the long-term effects that common management practices will have on the cycling and storage of key elements such as carbon (C), nitrogen (N) and phosphorus (P) (Conant et al. 2001, Soussana & Lemaire 2014, Fornara et al. 2016). Common management practices include animal grazing and the application to soils of different fertilizing materials including macro-nutrients (e.g. N, P, K, Mg) and lime (e.g. CaCO_3), all of which are known to influence grassland ecosystem functioning (Fornara et al. 2011, Isbell et al. 2013, Orwin et al. 2015, Sochorová et al. 2016). So far, a large number of research studies have focused on the relationships between management and changes in plant aboveground biomass, mainly because grassland productivity remains a key ecosystem service delivered to human society. Fewer studies have addressed potential long-term effects of different management practices (e.g. grazing, nutrient fertilisation and liming) on multiple plant belowground traits (e.g. total root mass, rates of root mass decomposition, root-microbial symbioses) in grassland ecosystems. Changes in root traits may have important consequences for soil nutrient cycling and the sequestration of C and N in grassland soils.

From an ecosystem functioning perspective, common grassland management practices greatly contribute to increase the availability of reactive inorganic forms of N and P in soils. Because of the close coupling of C and N cycles in soil ecosystems (Christopher and Lal 2007) changes in soil N availability may greatly affect soil C storage in human-managed grasslands (Lu et al. 2011, Riggs et al. 2015). Repeated nutrient additions can also influence total root mass and root C pools (Liu and Greaver 2010, Lu et al. 2011), mainly because of

the reduced role of roots in foraging for soil nutrients. For example, in natural or semi-natural ecosystems plant species tend to develop larger root systems, which maximise nutrient absorption especially in nutrient-poor soils (Hobbie 1992, Müller et al. 2000). Long-term repeated additions of multiple nutrients (i.e. NPKMg) to soils can significantly reduce plant biomass investment to belowground compartments (Fornara et al. 2013). This in turn will affect both the quantity and quality of root plant detritus returned to soils, thus potentially influencing rates of root mass decomposition.

Root litter quality (e.g. root N content, root lignin content, root C:N ratios), together with climatic factors and local soil conditions, plays a key role in regulating rates of root decomposition (Silver & Miya 2001, Fornara et al. 2009, Solly et al. 2014, Chen et al. 2017). Most root decomposition studies, however, have been carried out in natural and semi-natural ecosystems where human impact is generally lower. Thus, it is not clear how root decomposition might respond to multiple long-term grassland management practices.

Repeated nutrient fertilisation and animal grazing can also greatly influence root-microbial associations, such as root colonisation by arbuscular mycorrhizal fungi (AMF; Glomeromycota) (Hodge et al. 2001). Mycorrhizal fungi form a mutualistic relationship with roots, whereby fungi can provide soil nutrients to plants in return for the C supplied by host plants (Bardgett 2005). AMF are very common in grasslands (Johnson et al. 2003, Chapin III et al. 2012), however, their presence and abundance can be significantly affected by long-term nutrient fertilisation (Johnson et al. 2015, Liu et al. 2015, Propster and Johnson 2015, Sochorová et al. 2016). The common use of agricultural lime (non-nutrient fertilizer), which improves soil pH conditions might influence root decomposition processes through liming-induced increases in soil biological activity and/or changes in microbial composition (Fornara et al. 2011). Liming applications, for example, have been associated with higher AMF colonisation of roots (Johnson *et al.* 2005).

In this study we first address how (1) total root mass, (2) root mass decomposition, and (3) root AMF colonisation might respond to 22 years of grassland management, which include rabbit grazing, liming and chronic nutrient fertilisation. Secondly, we search for potential linkages between these root ‘traits’ and changes in important soil properties (i.e. soil N and P availability and total soil C, N and P pools). We hypothesize that plant belowground responses to long-term management will be mainly driven by the increased availability of organic and inorganic forms of nutrients (e.g. N and P), due to direct or indirect effects of grazing animals, artificial fertiliser and liming applications to soils.

Our study specifically tested the following hypotheses:

- (1) Total root mass will decrease whilst rates of root decomposition will increase under long-term multiple nutrient (e.g. NPKMg) additions;
- (2) Total root mass will decrease under grazing, partly because of increased nutrient additions to soils from grazing animals and partly because of plant aboveground responses to biomass removal by grazers (Holland et al. 1996);
- (3) AMF colonisation will increase in chronically N-fertilised soils, which have not received any inorganic P addition for decades thus leading to greater P-limitation for plant growth;
- (4) Liming-induced increases in soil pH by stimulating soil NO_3 availability (Kunhikrishnan et al. 2016) and microbial activity (Fornara et al. 2011) will lead to lower root mass, higher rates of root decomposition and greater AMF colonisation to acquire inorganic P from soils (Johnson et al. 2003).

We tested these hypotheses using data from a long-term grassland experiment where grazing, liming and five different nutrient treatments (i.e. Control, N-only, P-only, NPMg and NPKMg) have been applied for 22 years.

Methods

Study Site

We conducted our study at Silwood Park, Berkshire, UK, where a long-term grassland experiment (i.e. Nash's Field) was established in 1991 to test for the effects of common grassland management practices on plant species composition and plant yields (Crawley 1990; <http://www.imperial.ac.uk/silwood-park/research/silwood-lte/nashk-s/>). Nash's Field is a mesotrophic grassland described as *Lolio-Cynosuretum cristati* grassland, *Anthoxanthum odoratum* subcommunity according to the British National Vegetation Classification community (Rodwell 1992). This grassland is characterized by sandy, acidic soils of the Bagshot Series where introduced European rabbits (*Oryctolagus cuniculus*) have been particularly abundant since the 1950s. Before 1991 Nash's field was managed as a hay meadow and it is now dominated by perennial C3 grasses including *Agrostis capillaris*, *Festuca rubra* and *Holcus lanatus*. Common forbs include *Galium saxatile* and *Rumex acetosella*. A recent study of the Nash's Field shows how C3 grasses represent at least 73% of the total aboveground biomass in these experimental plots, whereas biomass of non-leguminous forbs varies between 10% and 16% and legume biomass represents <0.1% of the plant community biomass (Fornara et al. 2013).

Nash's Field is organized in a 4-factor split plot design (Appendix S1: Fig. S1). The experiment includes four large blocks (44 × 20 m each), two randomly selected to receive insecticide (– insects) and two with no insecticide applied (+ insects). Within each block the split-plot design incorporates two grazing plots (22 × 22 m each). Then, each grazing plot includes two liming plots (18 × 8 m each), and finally each liming plot incorporates five nutrient plots (2 × 2 m each). Thus, 4 blocks × 2 rabbit grazing treatments × 2 liming treatments × 5 nutrient treatments = 80 individual experimental units.

The insecticide treatment includes a cocktail of knock-down (Ambush, permethrin synthetic pyrethroid at 150 g a.i. ha⁻¹; Amvac Chemical, Surrey, UK) and systemic (Dimethoate-40; applied at 350 g a.i. ha⁻¹; BASF, Cheadle, UK) insecticides, which has been applied on three occasions each year (i.e. April, July, September) since 1992 (see also Allan and Crawley 2011). Rabbits were excluded from 1991, using fences constructed of wire mesh (3 cm diameter mesh size) attached to a rigid top wire, supported by wooden posts every 3.5 m (see Edwards and Crawley 1999). Each January, 5000 kg of CaCO₃ ha⁻¹ were applied to the limed plots. Each of the 80 small experimental plots (4 m²) has been managed to receive specific combinations of nutrient fertilisers (Crawley 1990). In this study, we specifically focused on the following 5 nutrient-treatments: (1) N-only, (2) P-only, (3) N + P + Mg, (4) all nutrients (N + P + K + Mg) and (5) no nutrients (i.e. control plots). Mineral nutrients are added annually to the soil and include 100 kg N ha⁻¹ in the form of ammonium nitrate (NH₄NO₃), 35 kg P ha⁻¹ in the form of triple superphosphate (Ca(H₂PO₄)₂), 225 kg K ha⁻¹ in the form of muriate of potash (KCl, K: 61% K₂O, Cl: 46%) and 11 kg Mg ha⁻¹ in the form of Epsom salts (MgSO₄).

Plant Belowground Sampling and Analyses

In May 2014, a 2.5 cm diameter soil corer was used to sample soils to a depth of 20 cm from five random locations within each of the 80 experimental plots included in our study. The five soil samples were homogenised and washed over a 2 mm size sieve with water to remove soil. Washed roots were oven-dried for 3 days at 60°C until they reached constant mass. Once dried, root samples were weighed to estimate total root mass (grams of dry roots per m²). A root subsample from each plot was ground to powder and analysed for total root C and N content following standard methods on a 1500 NA Carlo-Erba element analyser (Elan Tech, Lakewood, NJ, USA). Dry-root sub-samples were air-dried, ball milled to pass a 150

µm sieve and then processed for the determination of total P with the sodium hydroxide (NaOH) fusion method described by Smith and Bain (1982). A final root subsample was stored at 20°C for use in a root decomposition experiment (see below).

Soil Sampling and Analysis

A 2.5 cm diameter soil core (0-20 cm depth) was extracted in May 2014 from four random locations within each of the 80 experimental plots. Soils were sieved through a 2 mm size mesh and then dried for 3 days at 70°C. A subsample of the dried soil was analysed for total C (%) and N (%) content by combustion using a 1500 NA Carlo-Erba element analyser (Elan Tech., Lakewood, NJ, USA). Another subsample was used to estimate extractable phosphorus levels potentially available for plant uptake (i.e. Olsen P analysis) (Olsen et al. 1982). A final soil subsample was used to measure soil pH by creating a 1:5 soil:deionised water suspension.

Soil $\delta^{15}\text{N}$ natural abundance

Because all experimental treatments (i.e. grazing, liming and nutrient fertilization) either directly or indirectly tend to increase (and influence) soil N availability, we also measured the natural abundance of soil $\delta^{15}\text{N}$, which might help explaining potential long-term treatment effects on soil N cycling (Robinson 2001; Kleinebecker 2014). Soil subsamples collected in May 2014 from each experimental plot were sieved through a 2 mm mesh size. An 18 mg of this sieved soil was loaded into each of 80 individual tin capsules, which were then closed and weighed. Each capsule was placed in a well plate, which was then sent to the CHRONO laboratory at Queen's University Belfast, UK for $\delta^{15}\text{N}$ analyses. Soil samples were then dried and ground to powder and $\delta^{15}\text{N}$ of soil was determined using a stable isotope mass

spectrometer (Delta V™ Isotope Ratio Mass Spectrometer - Thermo Fisher Scientific, Waltham, MA USA).

Net Soil Nitrogen Mineralisation Rates

In June 2014 a 2.5 cm diameter soil corer was used to extract 3 randomly located soil samples to a depth of 20 cm in each of the 80 experimental plots. Samples were homogenised and sieved to a 2 mm mesh size, and stored for 12 hours in a cooler box before be processed in the laboratory. A 20 g subsample of soil from each plot was suspended in 50 ml of 1M KCl before being stored at 4°C overnight to allow the soil to settle. Once settled, a sample of the solution was transferred into tubes (8 ml) and analyzed for NH_4^+ and NO_3^- concentrations. A further 20 g soil sample from each plot was incubated at room temperature for 30 days.

Sufficient water was added to each sample after 2 weeks during the laboratory incubation to keep moisture constant. The same KCl extraction was then repeated after the 30 day-incubation period and final concentrations of NH_4^+ and NO_3^- were measured. Soil net N mineralization rates were calculated by subtracting initial from final NH_4^+ and NO_3^- concentrations (i.e. N mineralization rates were expressed per gram of soil).

Root Decomposition Experiment

We followed the protocol of Orwin et al. (2010) to estimate potential rates of root decomposition in a controlled laboratory incubation. A subsample of sieved soil (i.e. 31.05g of soil) was first added to each of 160 petri dishes (90 mm diameter) (2 petri dishes for each of the 80 experimental plots). Within each petri dish a 1 mm mesh size disc was placed on top of the soil, covering the entire surface area. Then 0.3 ± 0.05 grams of dried roots were placed on the fine mesh in each petri dish before being left to decompose in the dark at an ambient temperature of 25°C for 3 months. To test whether root decomposition might

respond more to (1) changes in local soil conditions under each experimental treatment or more to (2) changes in specific 'root properties' (i.e. root chemistry), we set up two parallel decomposition trials. In the first trial (including 80 petri dishes) a mixture of 'native' roots (i.e. 0.3 ± 0.05 g of dried root mass) from each plot (roots collected in May 2014) was decomposed in soils collected from the same plots (i.e. local roots decomposing on local soils). In the second trial we decomposed the roots of a 'non-native' plant species of Nash's Field, namely marram grass (*Ammophila arenaria*, which was collected in sand dune systems located along the north coast of Northern Ireland) under each of the same 80 local soil conditions used in the first trial (i.e. non-native 'control' roots decomposing on local soils). Roots were dried at 65°C for 7 days and a subsample was immediately ground and analysed for total C, N and P content. A sub-sample of the remaining dry roots (i.e. 0.3 ± 0.05 g) was then added to each of the 160 (i.e. 80 'control' marram grass and 80 'local' plant root mixtures) petri dishes (roots <2 mm diameter). All petri dishes were sealed (with the lid removed) using 75% duct tape and 25% parafilm, before being left in a dark room at an ambient temperature of 25°C between 26th June and 30th September 2014. Water was added to the petri dishes every two weeks to maintain an assumed soil moisture capacity of 9% by weight (Fornara et al. 2009) during the 3 months of the root incubation period. At the end of the incubation period roots were removed, dried for 7 days at 65°C, and weighed. Final root mass was compared to initial root mass to estimate the root mass loss (%) during the experiment. Dried root samples were then ground and analysed for total C, N and P (%).

Assessment of Mycorrhizal Infection

In July 2014 soil cores (2.5 cm diameter) were collected to a depth of 20 cm from 3 random locations within each of the 80 plots included in our study. Root samples were washed over a 2 mm mesh size and assessed for Arbuscular Mycorrhizal Fungi (AMF) colonisation using an

adapted version of the protocol developed by Vierheilig et al. (1998). Roots were cut into several 1 cm-long fragments and a random number of root fragments per plot were placed in a labelled cassette. A 90°C water bath was used to boil the cassettes in 10% KOH solution. After boiling for 10 minutes KOH solution was poured off and the cassettes were rinsed three times with tap water. Acetic acid was added until it covered all the cassettes and then poured off almost immediately and a 10% ink solution was finally added. The cassettes were boiled in the ink solution for 5 minutes, then the ink solution was poured off and the cassettes were rinsed 3 more times with tap water before being added to lactoglycerol and left to destain overnight. Roots were mounted in lactoglycerol on labelled microscope slides, running parallel to each other, before being analysed using an Olympus BH2 microscope. Each slide was analysed at magnification x 400 and a crosshair was placed in the eyepiece. An adaptation of the protocol developed by McGonigle et al. (1990) was used for the identification of AMF structures present in the roots, namely hyphae, arbuscules and vesicles were identified and recorded. A total of 100 stops was the aim per slide and a tally of structures that intersected with the crosshair was taken per stop.

Data Analysis

We used a mixed effects ANOVA model where the four blocks were included as random factors and the experimental treatments (i.e. insecticide, grazing, liming and nutrient applications) represented our (fixed-effect) predictor variables (Appendix S1: Fig. S1). Since the pesticide treatment did not account significantly for variation in any of our response variables, we then produced simpler models, which included only grazing, liming and nutrient as fixed effects. Grazing, liming and nutrient fertilisation treatments are nested within each other and then replicated across four randomly located larger plots (i.e. blocks).

We tested for potential effects of grazing, liming and nutrient applications (and their interactions) on multiple soil and plant belowground parameters (e.g. total root biomass, root mass decomposability, nutrient accumulation indexes, AMF colonisation etc.). When necessary values were square root transformed to improve normal distribution of residuals. Restricted Maximum Likelihood (REML) method was used to produce final models, which were checked to ensure they conformed to modelling assumptions. Significant differences between factor levels were tested using post-hoc Tukey tests. We also performed simple and multiple regression analyses to identify potential relationships between plant and soil variables, with particular focus on the correlation these variables with soil pH.

We calculated the percent root mass loss as an indicator of root decomposability using the following formula:

$$\left(1 - \frac{\text{final root mass}}{\text{initial root mass}}\right) \times 100$$

We calculated the Nutrient Accumulation Index (NAI) as an index of nutrient dynamics in decomposing root material (Fornara et al. 2009). We distinguished between NAI (for Nitrogen) and PAI (for Phosphorus). Thus, NAI or PAI is equal to:

$$\frac{W_t \times X_t}{W_o \times X_o}$$

Where W_t = root dry mass remaining at time t , W_o = initial root dry mass, X_t = root nutrient concentration at time t , X_o = initial root nutrient concentration. We assumed that NAI = 1 would suggest no changes in mass balance for nutrient X (either N or P), $NAI < 1$ would suggest net mineralization of nutrient X , and $NAI > 1$ would suggest net immobilization of nutrient X .

To estimate the degree of AMF colonisation of plant roots we first counted AMF structures (arbuscules, hyphae and vesicles) present on between 14 and 25 root fragments on each microscope slide. We then calculated the proportion of each structure as:

$$\frac{\text{Count of structure per slide}}{\text{Total number of stops per slide}} \times 100$$

Total AMF colonisation per slide was estimated by adding the proportion of all structures together (see McGonigle et al. 1990), however, the relative abundance of arbuscules per slide was used in our data analysis due to their presence giving a better indication of mutualistic functioning (Johnson et al. 2003).

Results

Total root biomass

We found that grazing did not have any significant effect on total root biomass ($F_{1,60} = 0.7$; $P = 0.35$; Fig. 1a), while liming was associated with a significant decrease in root mass ($F_{1,60} = 15.9$; $P = 0.0002$; Fig. 1b). Total root mass in limed plots was 30% lower than in un-limed plots. Nutrient additions also had a significant negative effect on total root mass ($F_{1,60} = 3.1$; $P = 0.016$), which was mainly determined by root mass decreases associated with the All (i.e. NPKMg; Tukey's *posthoc* test, $P < 0.05$) nutrient treatment (Fig. 1c).

Root mass decomposition

Grazing did not affect root mass loss (%) of either native (from Nash's Field) ($F_{1,60} = 0.02$, $P = 0.88$) or control (marram grass) roots ($F_{1,60} = 0.37$; $P = 0.55$; Fig. 2a). The addition of lime significantly increased root mass loss of both control (+8% for the marram grass) ($F_{1,60} = 18.2$, $P < 0.0001$) and native root species (+4%) ($F_{1,58} = 5.84$; $P = 0.02$; Fig. 2b). Nutrient additions did not have any significant effect on root mass loss (%) of either native plant species (from Nash's Field) ($F_{4,60} = 0.34$, $P = 0.85$) or the (non-native)-control marram grass ($F_{4,60} = 0.99$, $P = 0.42$; Fig. 2c).

Root nutrient accumulation indexes

Grazing did not affect the nitrogen accumulation index (NAI) of native ($F_{1,60} = 0.55$; $P = 0.46$) or control roots ($F_{1,60} = 0.30$, $P = 0.59$; Fig. 3a). Also, grazing did not affect the Phosphorus Accumulation Index (PAI) of native ($F_{1,60} = 0.45$, $P = 0.51$) or control roots ($F_{1,60} = 0.030$, $P = 0.86$; Fig. 4a). The addition of agricultural lime caused a significant decrease in NAI of control ($F_{1,60} = 4.8$, $P = 0.03$) and native roots ($F_{1,59} = 5.72$; $P = 0.02$; Fig. 3b). Liming did not affect PAI of native ($F_{1,60} = 0.09$; $P = 0.76$) or control roots ($F_{1,60} = 3.88$; $P = 0.05$; Fig. 4b), despite the trend observed suggesting a negative effect of liming.

Nutrient addition did not affect NAI of either native ($F_{4,60} = 2.26$; $P = 0.07$) or control roots ($F_{4,60} = 0.73$, $P = 0.58$; Fig. 3c). PAI of native roots was not significantly affected by nutrient addition ($F_{4,60} = 0.51$; $P = 0.73$; Fig. 4c), while PAI of control roots was ($F_{4,60} = 8.57$; $P < 0.0001$).

Root AMF colonisation (%)

We found no significant effects of grazing on total AMF colonisation (%) of plant roots in the Nash's Field ($F_{1,60} = 3.15$, $P = 0.15$; Fig. 5a; Appendix S1: Table S1), whereas the addition of lime caused a significant increase (+22% compared to unlimed soils) in total AMF colonization ($P = 0.001$; Fig. 5b; Appendix S1: Table S1). We also found that nutrient additions caused significant changes ($P = 0.003$) in total AMF colonisation, whereby N-only additions were associated with greater AMF abundance whereas the addition of multiple nutrients (i.e. NPKMg or NPMg treatments) was related to lower AMF colonisation (Fig. 5c; Appendix S1: Table S1). Grazing did not significantly affect other AMF structures while the addition of agricultural lime and nutrients influenced arbuscule and hyphae abundance but not vesicule abundance (Appendix S1: Table S1).

Soil pH and belowground properties

Soil $\delta^{15}\text{N}$ natural abundance was not significantly affected by grazing or nutrient additions but it was significantly negatively affected by repeated liming applications ($F_{1,60} = 13.5$, $P < 0.0001$). Liming also increased soil nitrate levels (Appendix S1: Fig. S2) and Nitrogen Use Efficiency (NUE; Appendix S1: Fig. S3). NUE was calculated as the ratio of plant yields (i.e. total aboveground plant biomass; g m^{-2}) and total N uptake by the plant community (g N m^{-2}).

We found that most soil parameters were significantly affected by the liming treatment and that the liming effect was due to strong liming-positive effects ($F_{1,79} = 532$, $P < 0.0001$) on soil pH (see also Appendix S1: Fig. S4). For example, soil pH was significantly negatively related to total root mass ($F_{1,79} = 7.4$, $P = 0.008$; Fig. 6a), total AMF (%) colonisation ($F_{1,79} = 12$, $P = 0.001$), NAI of control roots ($F_{1,79} = 4.6$, $P = 0.03$) and to the decomposition of both native (Fig. 6b) and control roots (Fig. 6c). Soil pH was significantly positively related to soil NO_3 concentrations ($F_{1,79} = 12.2$, $P = 0.0008$; Fig. 7a) and negatively related to soil $\delta^{15}\text{N}$ ($F_{1,79} = 20.8$, $P < 0.0001$; Fig. 7b). However, multiple regression analyses show that soil pH and soil C and N stocks were not significantly related (Fig. 7c, d).

Discussion

Overall, our results show how the common practice of agricultural liming (i.e. the addition of CaCO_3 to soils) strongly affected total root mass, root decomposition, root AMF colonisation as well as soil N availability and cycling. Our evidence is that liming effects were mainly driven by liming-induced increases in soil pH, which influenced multiple root and soil parameters more than animal grazing or chronic nutrient fertilisation.

According to our first hypothesis we found that total root mass was significantly reduced by the long-term addition of multiple nutrients (i.e. NPKMg). This result could be associated with the positive effects that repeated fertilizer applications have on soil nutrient

availability. Increased soil nutrient availability determines shifts in plant resource allocation from belowground to aboveground plant compartments thus causing a reduction in total root mass (Hobbie 1992, Müller et al. 2000, Hobbie et al. 2010, Zeng et al. 2015). Reductions in total root mass were, however, lower under the N-only and P-only treatments (Fig. 1c), possibly because plants still need to invest in ‘foraging’ root mass to acquire nutrients, which are less available in soils. Contrary to expectations, we did not find any clear evidence that different nutrient fertilisation applications influenced the decomposition of either control or native roots. Our results show that the addition of all nutrients together (i.e. NPKMg treatment) was associated with increases in native root decomposition whereas the N-only treatment was associated with decreases in native root decomposition but both these effects were not statistically significant (Fig. 2c).

Contrary to our second hypothesis we found no significant grazing effects on total root mass. This is despite the expectation that animal grazing would cause a shift in plant resource allocation from belowground to aboveground compartments to replace the photosynthetic tissue removed by herbivores (Holland and Detling 1990). The lack of significant grazing effects on root mass may be related to shifts in plant species composition under repeated herbivore pressure whereby unpalatable plants are able to maintain their root systems while palatable plants reallocate resources to replace lost aboveground biomass (Schuster 1964). In these grasslands, for example, it was shown that *Cirsium arvense* (a perennial broadleaf plant species with extensive root systems) greatly benefits from animal grazing mainly because of competitive release from other plant species, which are selectively removed by grazers (Edwards et al. 2000). Net grazing effects on plant belowground mass will ultimately depend on how grazer ‘identity’ (i.e. rabbits vs. ungulates vs. arthropods) interacts with resistance/resilient traits of dominant plant species to influence soil nutrient availability.

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According to our third hypothesis, we found that root AMF colonisation was significantly lower under chronic multi-nutrient applications (i.e. NPKMg and NPMg), which increased soil nutrient availability including inorganic forms of P. As opposite, AMF abundance was highest under the N-only treatment, a finding, which may further confirm the key role of AMF in P acquisition (Wang et al. 1993, Guo et al. 2012). In fact, it is likely that long-term N-only fertilized soils have become 'P-limited' with relatively high soil N:P ratios (see Appendix S1: Fig. S5) a N-to-P stoichiometric imbalance, which has likely promoted AMF colonization of these roots. AMF colonisation was not significantly influenced by grazing despite previous findings showing that AMF increases colonisation under grazing (Jansa et al. 2006, Wearn & Gange 2007).

According to our fourth hypothesis we found that liming had strong positive effects on soil N availability (e.g. soil NO₃ levels) and on soil N cycling and that liming-induced increases in soil pH significantly influenced both belowground and aboveground compartments. Increases in soil pH have previously been shown to enhance soil microbial activity and soil N availability (Van Aarle et al. 2002, Kemmitt et al. 2006, Rousk et al. 2011), with significant implications for plant N uptake. For example, we suggest that decreases in soil $\delta^{15}\text{N}$ natural abundance under increasing soil pH values (Fig. 7b), bring evidence of more efficient N uptake by plants in limed soils. This is supported by our finding that Nitrogen Use Efficiency (NUE) in limed plant communities was higher than in un-limed plants (Appendix S1: Fig. S3). These results also confirm those from a recent study (Kleinebecker et al. 2014), which show how soil $\delta^{15}\text{N}$ signals decreased under greater plant species diversity because of a more efficient use of N in species-rich grassland communities. Thus both increased soil nitrate levels and more efficient plant N uptake may explain the reduced need for plants to invest in large root systems in repeatedly limed soils (Fig. 1b). It is known that agricultural liming tends to improve soil cation exchange capacity, ultimately

increasing aboveground productivity (Johnson et al. 2005; Fornara et al. 2011) rather than root productivity. Our results also show that liming was associated with higher root decomposition, which could be driven by liming-induced increases in nutrient availability and cycling (Rangel-Castro et al. 2005, Mijangos et al. 2010, Wachendorf 2015). Our evidence is that liming contributed to increase the mineralisation of N in decomposing root material (Fig. 3a; NAI decreased under lime addition), suggesting that microbes could satisfy their metabolic N requirements faster in limed soils than in un-limed soils. Liming, however, did not significantly influence changes in P mineralisation-immobilisation in root decomposing material (Fig. 4a), which was significantly influenced by nutrient fertilisation (at least in the control root samples). Control roots decomposing on soils conditioned by the N-only treatment showed a trend toward higher P mineralisation (i.e. low PAI, Fig. 4c) perhaps due to P limitation compared to N in these soils. We found, however, that liming was associated with increases in AMF colonisation, and this is possibly due to increased P immobilisation in limed soils, which could lead to lower P availability for plant uptake. Higher soil pH values are usually associated with reduced P availability in soils due to P binding with the calcium (Ca) component of lime to form phosphate compounds (Aerts et al. 2003, Olsson et al. 2009, Saarsalmi et al. 2011).

Despite our expectation that grazing would enhance nutrient cycling and hence decrease NAI, our results do not confirm this and differ from previous studies which found N mineralisation-immobilisation dynamics to be directly affected by grazing (Shariff et al. 1994, McNaughton et al. 1998, Coetsee et al. 2010). It could be that grazing interactions with nutrient fertilisation and liming have reduced net grazing effects on soil nutrient cycling. A recent study from natural tundra ecosystems show how nutrient fertilisation increased soil N availability in ungrazed rather than grazed plots, and how liming could partly negate nutrient fertilisation effects on soil N availability (Stark et al. 2012).

Based on positive correlations between root decomposition and soil pH we would have expected soil C and N pools decrease in limed soils. However, we did not find this evidence in our study, where limed and unlimed plots had similar C (%) content and similar soil C stocks. Previous studies also found weak correlations between soil pH and soil C and N stocks (Kemmitt et al. 2006), and this could be related to the key role of microbial communities and their activities. Recent studies clearly show positive effects of liming on soil C stocks across different ecosystems (Fornara et al. 2011, Melvin et al. 2013), while findings from a review study (Paradelo et al. 2015) suggest that liming can cause both increases and decreases in soil C stocks. Liming-induced increases in the activity of microbes and fauna within the soil would increase the transformation of organic matter via decomposition and mineralisation thus increasing C losses from soils. On the other hand, liming induced changes in soil structure (as Ca in lime binds with soil ions) would lead to stabilization of soil macro-aggregates and thus to higher soil C accumulation (Kunhikrishnan et al. 2016). It could also be that higher microbial activities in limed soils will contribute to increase microbial waste and thus C detritus, which is incorporated and stabilized into smaller soil organo-mineral fractions (Fornara et al. 2011).

Conclusion

Our results bring evidence of how liming applications can strongly influence multiple soil and plant properties (and traits) including soil N cycling, total root mass, root decomposability and root AMF colonisation. Liming-induced increase in soil pH is a key factor explaining these belowground responses, however, changes in soil pH were not related to changes in soil C and N stocks in this permanent grassland. These findings suggest that regular liming applications could contribute to maintaining grass yields while not having detrimental effects on the C sequestration ability of grassland soils. Moreover, the addition of

agricultural lime could significantly reduce the need for artificial nutrient fertilizers because of liming positive effects on soil nitrate levels and on root AMF colonization (which improves P uptake by plants). We suggest that regular liming applications together with reduced artificial nutrient fertilization could contribute to maintaining grass yields without compromising the delivery of other ecosystem services such as soil nutrient cycling and soil C sequestration, thus enhancing the long-term sustainability of managed grasslands.

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Contribution:

J. Heyburn and D. Fornara equally contributed to write the manuscript. All authors significantly contributed to the research study.

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Supporting Information

Additional supporting information may be found in the online version of this article at

<http://onlinelibrary.wiley.com/doi/10.1002/eap.xxxx/supinfo>

Data Availability

Data available from the Dryad Digital Repository: <http://dx.doi.org/10.5061/dryad.p0s23>

Fig. 1. Effects of animal grazing (a), liming (b) and nutrient additions (c) on total root mass.

YES = + treatment; NO = – treatment. Error bars (SE) indicate variation in root mass between plots that received the same treatment. Different letters indicate a post-hoc difference in means at $p < 0.05$, NS = Not significant; $p \geq 0.05$.

Fig. 2. Effects of grazing (a), liming (b) and nutrient additions (c) on root mass loss (%) from either ‘native’ or ‘control’ roots. ‘Native’ = mix of dry roots collected from each experimental plot in the Nash’s Field. ‘Control’ = dry roots of a non-native control grass species (i.e. *Ammophila arenaria*). Error bars (SE) indicate variation in root mass between plots that received the same treatment. Different letters indicate a post-hoc difference in means at $p < 0.05$, NS = Not significant; $p \geq 0.05$.

Fig. 3. Effects of animal grazing (a), liming (b) and nutrient additions (c) on mean “Nitrogen Accumulation Index” (NAI). Error bars (SE) indicate variation in between plots that received the same treatment. Different letters indicate a significant difference, NS = Not significant.

Fig. 4. Effects of animal grazing (a), liming (b) and nutrient additions (c) on mean “Phosphorus Accumulation Index” (PAI). Error bars (SE) indicate variation in PAI between plots that received the same treatment. Different letters indicate a significant difference, NS = Not significant.

Fig. 5. Effects of animal grazing (a), liming (b) and nutrient additions (c) on total AMF colonisation expressed as percent of total roots infected. Error bars (SE) indicate variation in AMF colonisation between plots that received the same treatment. Different letters indicate a significant difference, NS = Not significant.

Fig. 6. Relationships between soil pH and multiple root traits, which include total root mass (g m^{-2}), native root mass loss (%), and control root mass loss (%). Shaded area refers to 95% Confidence Intervals.

Fig. 7. Relationships between soil pH and key soil parameters, which include soil NO_3^- concentration (mg kg^{-1}), soil $\delta^{15}\text{N}$, soil N and soil C stocks (t ha^{-1}). Shaded area refers to 95% Confidence Intervals.