Plant-soil feedback is shut down when nutrients come to town

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Abstract

Background and aims The concept of plant-soil feedback is increasingly used to explain plant community assembly processes. Soil nutrient availability can be expected to play a critical role on these processes. However, little is known about the effects of nutrient availability on feedback direction and strength.

Methods A plant-soil feedback experiment was performed with the grasses Anthoxanthum odoratum and Festuca rubra, and the forbs Leontodon hispidus and Plantago lanceolata, on soil with either low or high nutrient availability. Additionally, we tested if plant-soil feedback of the two forbs under these conditions changed by inoculation of the soil with spores of an arbuscular mycorrhizal fungus.

Results Increased nutrient availability neutralised plant-soil feedback based on shoot biomass independent of its negative or positive direction, whereas the effects on root biomass were either not altered or turned negative. Mycorrhizal fungi spore addition decreased negative feedback and increased positive feedback.

Conclusions Our results suggest that negative plant-soil feedback on low nutrient soil can be overcome with nutrient addition, and that positive soil biota associations on low nutrient soil may become superfluous with nutrient increase. We hypothesize that species-specific, microbial mediated plant community assembly processes occur in low rather than high nutrient environments.

Keywords Plant-soil feedback · Soil nutrients · Mycorrhizae · Soil biota · Species-specific · Plant community assembly · Anthoxanthum odoratum · Festuca rubra · Leontodon hispidus · Plantago lanceolata

Introduction

Plant species co-existence, succession and invasion processes are commonly linked to responses between the plant and its surrounding soil. This so called plant-soil feedback is defined as changes to soil properties induced by plants, which in turn affect these same or other individuals in the plant community positively or negatively (van der Putten et al. 2013). The term plant-soil feedback thus encases a large variety of soil property changes; however, it is particularly the microbial changes that are considered to play driving roles in plant community assembly (e.g., Van der Putten et al. 2013; Bever et al. 2015). Key in this theoretical framework is the accumulation of species-specific microbial communities in and around the plant’s root system, which generate species-specific plant-soil feedback ranging...
from strongly negative to moderately positive (e.g., Klironomos 2002; Petermann et al. 2008; Cortois et al. 2016). Especially negative feedback is regarded as imperative to plant species community processes, because negative feedback is a form of negative frequency dependence implying that successful species limit their own growth when in high abundance (e.g., Bever et al. 1997, 2015). There has been considerable progress in understanding the processes that link plant-soil feedback to such natural plant community assembly processes. However, how the environment affects plant-soil feedback strength and direction has yet received relatively little attention (Ke and Miki 2015; Lekberg et al. 2018; Erktan et al. 2018).

Belowground communities are suggested to be predominantly structured by environmental conditions such as soil fertility (Ramirez et al. 2010; Tedersoo et al. 2014; Lekberg and Waller 2016; Laliberté et al. 2017). Increases in soil nutrient availability decreases soil microbial biomass (Treseder 2008; Mitchell et al. 2010; Ramirez et al. 2012, but see Manning et al. 2008), and shifts bacterial, fungal and archaeal soil community composition (e.g., Ramirez et al. 2010; Bates et al. 2011; Leff et al. 2015). These shifts in microbial community composition were consistent in grasslands across the globe (Leff et al. 2015), and, for nitrogen enrichment, also across biomes in general (Fierer et al. 2009; Ramirez et al. 2012). Such consistent changes are expected to interact with plant growth and hence to affect plant-soil feedback driven community assembly processes (e.g., Ehrenfeld et al. 2005; Luo et al. 2017). However, the few studies testing nutrient availability yield mixed results (Manning et al. 2008; Petermann et al. 2008; Harrison and Bardgett 2010), introducing two contrasting theories on how nutrient addition affects plant-soil feedback strength and direction via microbial community composition shifts.

Plant-soil feedback is, on the one hand, thought to become increasingly negative with increasing nutrient availability, because nutrients are expected to stimulate antagonistic soil biota and suppress protagonists (Rúa and Umbanhowar 2015; Revillini et al. 2016; van der Putten et al. 2016; Lekberg et al. 2018). This complies with studies showing an increase in pathogen abundance in the soil with increasing nutrient availability (Paungfoo-Lonhienne et al. 2015; Wei et al. 2018; Veach et al. 2018). Moreover, this increase in soil pathogens was shown to result in increased negative plant-soil feedback of Panax notoginseng, Chinese ginseng, with high nitrogen addition to this production system (Wei et al. 2018). However, for more natural systems, this remains untested.

Soil protagonist studies mainly focus on widespread mutualistic interactions between plants and mycorrhizal fungi that provide a competitive advantage to the plant via nutrient uptake and protection against pathogens (Sikes 2010). Stimulation of the mycorrhizal fungi community resulted largely in positive plant-soil feedback effects (Anacker et al. 2014; García-Parisi and Omacini 2017). These interactions generally decrease with nutrient addition (e.g., Egerton-Warburton and Allen 2000; Corkidi et al. 2002; Leff et al. 2015) suggesting that suppression of mycorrhizal protagonists via soil nutrient increase, strengthens negative feedback. However, we are unaware of any studies that have tested this interaction effect between nutrients and protagonist soil biota on plant-soil feedback.

On the other hand, theory predicts plant-soil feedback to become less negative with increasing nutrient availability. This may result from decreased nutrient stress, increase in overall fitness and an elevated tolerance to soil-borne pathogens by the plants in high nutrient environments. Support for this opposing hypothesis can be found in experiments by Petermann et al. (2008), who showed a partial decrease in negative plant-soil feedback across 24 European grassland species with nutrient addition. This was tested by comparing plant growth between soils where the same plant species had grown on before (own conditioned soil) and soil where another species had grown on (foreign conditioned soil) with and without nutrient addition. However, the authors note that the observed partial decrease in negative plant-soil feedback with nutrient addition could result from either an interaction effect of nutrients and soil biota, or from reducing nutrient limitation on own soils conditioned by acquisitive species compared to foreign conditioned soils of more conservative species.

This overview shows that the literature on interactions between plant soil-feedback and soil nutrients is far from conclusive. Here, we tested this interaction and, moreover, assessed the role of common soil protagonists in plant-soil feedback relations. In particular, we performed a greenhouse plant-soil feedback experiment with two grasses, Anthoxanthum odoratum and Festuca rubra, and two forbs, Leontodon hispidus and Plantago lanceolata, on low and high nutrient soil. For the species that showed a fast response to mycorrhizal fungi spore
addition in the seedling phase, Leontodon and Plantago, we stimulated mycorrhizal root colonisation by adding mycorrhizal spores to the seedlings in a full-factorial design with soil nutrient availability. We asked (i) whether nutrient addition changes plant-soil feedback direction and strength, and (ii) whether increased exposure to arbuscular mycorrhizal fungi affects plant-soil feedback and how this response interacts with the response to nutrient availability. Based on our results, we discuss the role of nutrients and mutualistic interactions in shaping plant-soil feedback and the implications of these factors on plant community assembly processes.

**Material and methods**

**Experimental design**

A plant-soil feedback pot experiment with a conditioning and a feedback phase was performed for two grasses, Anthoxanthum odoratum and Festuca rubra, and two forbs, Leontodon hispidus and Plantago lanceolata. In the feedback phase, each species was grown on soil that had been conditioned by the same species, and, as a control, on a mixture of all four conditioned soils. All species were also grown with low and high nutrient availability in a full-factorial design with soil history \( (n = 6; \text{Fig. 1}) \). A pilot experiment showed that addition of spores of the generalist *Rhizophagus irregularis* mycorrhizal fungus to seedlings (details see below) significantly increased mycorrhizal fungi colonisation in roots of Leontodon and Plantago, but not in roots of Anthoxanthum and Festuca (Fig. S1). Shoot biomass was not significantly affected, indicating that inoculation with mycorrhizal spores did not have immediate negative, parasitic effects on plant growth (data not shown). Only Leontodon and Plantago were, therefore, considered to show a sufficiently rapid response to mycorrhizal spore addition and, as a result, to potentially be affected in their plant-soil feedback by stimulation of mycorrhizal fungi colonisation. These two species were in the feedback phase also grown with and without addition of spores of the mycorrhizal fungus in a full-factorial design with nutrient status and soil history \( (n = 6; \text{Fig. 1}) \).

**Conditioning phase**

Seeds of the four species (De Bolderik, Wervershoof, the Netherlands) were surface sterilised by shaking in 1.35% commercial bleach (diluted 1:1) for 30 min at 140 rpm. After thorough rinsing with demi-water, seeds were incubated in sterile petri-dishes (ø 9 cm) lined with wet filter paper. Growth room conditions for germination were 25 °C during the 12 h light period (MASTER TL-D Reflex 18 W/840 lights, Phillips Lighting B.V., Eindhoven, the Netherlands) at, on average, 55 μmol

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**Fig. 1** Experimental setup of the feedback phase. Four species, Anthoxanthum odoratum, Festuca rubra, Leontodon hispidus and Plantago lanceolata, were grown on soil conditioned by the same species (own) and on a mixture of all four conditioned soils (mix). Both own and mix soil received no addition of nutrients (low) and addition of a nutrient solution each week (high). Leontodon and Plantago were, in addition, planted without (−) and with addition of spores of a mycorrhizal fungus (+). This was done in a full-factorial design and resulted in 24 treatments \( (n = 6) \).
PAR m$^{-2}$ s$^{-1}$ at petri dish level and 10 °C during the 12 h dark period. Seedlings were carefully transferred to pots once their cotyledon(s) had emerged. Seed germination of the different species was timed to have the same transfer day.

Eight seedlings per species were transferred into disinfected (3 d in 1% MENNO clean, KaRo BV, Zwaagdrijk, the Netherlands), free draining 1.5 L pots (top diameter 13 cm, bottom diameter 11.5 cm, height 15 cm) lined with root cloth at the bottom ($n=48$). Pots were filled with a loamy soil (30% sand, 60% loam, 10% clay; Schinnen, the Netherlands) sieved at 1 cm mesh and placed in dishes to prevent nutrient leaching. Seedlings were thinned to four evenly distributed individuals per pot within the first 10 d and grown for 8 weeks in the greenhouse between 20 and 24 °C during the 16 h light period (MASTER GreenPower Plus 1000 W EL/5X6CT lights, Phillips Lighting B.V., Eindhoven, the Netherlands) at minimally 250 μmol PAR m$^{-2}$ s$^{-1}$ at pot level and 17–20 °C during the 8 h dark period. Pots were watered 3–5 times a week with deionised water. After 8 weeks, the shoot was cut off below the growing point and soil with roots was cut into clumps of 2–3 cm. All cut soil pots of each species were pooled and thoroughly mixed, and the mix of the four species-specific soils was made by pooling and thoroughly mixing half of the total amount of each conditioned soil (w/w). For each conditioned soil, plant available nitrogen (N) and potassium (K) (mg kg$^{-1}$ dry soil) was determined by adding 50 mL of 0.2 M NaCl to 10 g of fresh soil, shaking for 2 h at 120 rpm and filtering the soil out. NO$_3^-$, NH$_4^+$ and K$^+$ concentrations were measured on an AutoAnalyzer 3 (Bran+Luebbe GmbH, Norderstedt, Germany) ($n=3$). Plant available phosphorous (P) (mg kg$^{-1}$ dry soil) was determined according to Olsen et al. (1954) and measured on an ICP-OES iCAP 6000 (Thermo Fisher Scientific Inc., Waltham, MA, USA) ($n=3$). Mainly NO$_3^-$ levels between the own soil of the four species and the mix soil differed at the start of the feedback phase (Fig. S2). The amount of NO$_3^-$ added over the whole growing period (7480 μmol kg$^{-1}$) in the high nutrient treatment was 4 to 7 fold larger than the difference in NO$_3^-$ that occurred between own and mix soil treatments. Initial differences in nutrient concentrations in the conditioned soils were thus minor compared to nutrient addition treatments.

Feedback phase

From a second batch of seedlings that were germinated in the same way as for the conditioning phase, eight seedlings per species were transferred into the same size of pots as in the conditioning phase and thinned to four individuals per pot within the first 10 d. Leontodon and Plantago seedlings with mycorrhizal fungi addition were planted in dents filled with a mixture of 2.9 g dry soil with 0.1 g Rhizophagus irregularis spores in attapulgit powder (approximately 1000 spores per dent, 4000 per pot; strain DAOM197198; Symplanta GmbH & Co. KG, Darmstadt, Germany). Pots were watered every 3–5 d with deionised water. Once a week, with exception of week six, high nutrient pots received 200 mL nutrient solution (9350 μmol L$^{-1}$ NO$_3^-$, 1115 μmol L$^{-1}$ NH$_4^+$, 4686 μmol L$^{-1}$ K$^+$ and 930 μmol L$^{-1}$ PO$_4^{3-}$) instead of deionised water. Nutrients in the high nutrient treatments had been largely depleted to comparable levels as in the low nutrient treatments at harvest (Fig. S3). Pots were placed in pairs per species, own and mix soil, in a randomised block design in the greenhouse and grown for 6 weeks under comparable conditions as in the conditioning phase. Leaf counts indicated that initial positive soil history effects occurring for some species in the first 2 weeks faded or turned negative over time consistent with Dudenhöffer et al. (2018) (data not shown). This indicates that the time period of our feedback phase was long enough to observe an accumulation of biotic soil effects. After 6 weeks, shoots including flowers were cut off at the base, dried at 70 °C for at least 72 h and weighed. Roots were washed out of the soil on a sieve with tap water during which obvious dead roots from the previous phase were discarded. Approximately 20 mg (fresh weight) of these cleaned roots were stored in 70% ethanol to determine mycorrhizal colonisation. The remainder was dried at 70 °C for at least 72 h and weighed.

Mycorrhizal fungi staining and counting

Roots stored in 70% ethanol were cleared in 10% KOH (w/v) for 1 d, after which the solution was refreshed and roots were stored for another 1–2 d, until roots were sufficiently cleared (duration was species dependent). After this, roots were rinsed with 85% ethanol until no pigment leached out anymore. Roots were then placed in 5% HCl for 5 min at room temperature. Mycorrhizal
fungi were stained by placing roots in a 0.05% trypan blue solution (10 mL glacial acetic acid, 200 mL glycerol, 0.2 g trypan blue and 190 mL dH2O) at room temperature for 1–2 h (duration was species dependent). Roots were stored in acid glycerol overnight and stored in fresh acid glycerol until counting. Approximately ten, randomly selected, cleared and stained root pieces of approximately 1 cm were randomly placed on a microscopic slide with a 0.1 mm grid. Horizontal grid lines were traced under a light microscope and on the first 100 grid intersects with a root fragment, mycorrhizal structures (hyphae, vesicles and spores; arbuscules were not observed) were noted according to McGonigle et al. (1990). Relative mycorrhizal colonisation was calculated by dividing the number of incidences that a mycorrhizal structure was observed by the number of incidences that no mycorrhizal structure was observed ($n = 3$).

Calculations and statistics

Plant-soil feedback was calculated on the average biomass of the four individuals in a pot and for each pair of own and mix treated soils as (Dry weight$_{own}$ soil $-$ Dry weight$_{mix}$ soil) / Dry weight$_{mix}$ soil, both based on either shoot or root dry weight ($n = 6$). We analysed all data using ANOVA with generalised least square (gls) models from the nlme package (Pinheiro et al. 2017) in R 3.4.3 (R Core Team 2017). Model validation followed recommendations of Zuur et al. (2010). Full models included species, nutrient and, in case of biomasses and soil nutrients, soil history, and their interactions as fixed factors. Additionally, models testing the effect of mycorrhizal spore addition included the fixed factor mycorrhizae addition with all its interactions.

Inclusion of the factor species often resulted in heterogeneity of variances. In these cases, models were run with a variance structure allowing for differences in variance between species according to Zuur et al. (2009), using varIdent from the nlme package (Pinheiro et al. 2017). Full models including mycorrhizae addition often showed heterogeneity of variances because of the factor nutrient and, in some cases, mycorrhizae addition, and were therefore run with a weighted variance structure of one or two of these factors (Zuur et al. 2009). Full models for Leontodon and Plantago including mycorrhizae addition and also soil history as fixed factor were run per species, because heterogeneity of variances could not be overcome due to low statistical power. Data on NO$_3^-$ in the feedback phase was ln-transformed to meet assumptions of a normal distribution of the model residuals. After the full models were defined, post-hoc tests were run on these models using the emmeans-package (Lenth 2018).

Results

Nutrient addition eliminated plant-soil feedback

Plant-soil feedback of two grasses, Anthoxanthum and Festuca, and two forbs, Leontodon and Plantago, was determined on soil with low and high nutrient availability. Nutrient addition resulted in higher aboveground biomass of every species in each soil, whereas belowground biomass was not always affected by nutrient addition (Fig. S4). For plant-soil feedback based on shoot biomass, no overall effect of plant species or nutrient addition occurred. However, a significant interaction between plant species and nutrient addition was observed (Table 1). Plant-soil feedback based on shoot biomass of the two grasses Anthoxanthum and Festuca was absent and not significantly affected by nutrient addition (Fig. 2a). Leontodon and Plantago, on the other hand, showed positive and negative shoot feedback on low nutrient soil, respectively. Addition of nutrients largely eliminated this feedback (Fig. 2a).

For plant-soil feedback based on root biomass, no overall significant effect of nutrient addition occurred (Table 2). Plant-soil feedback based on root biomass was negative for Anthoxanthum and neutral for Festuca and Plantago independently of the nutrient addition treatment (Fig. 2b). Leontodon, on the other hand, showed neutral feedback on low nutrient soil, which turned negative with addition of nutrients (Fig. 2b).

| Table 1 ANOVA results of species, nutrient addition and their interaction on plant-soil feedback based on shoot and root biomass |
|---|---|---|---|---|---|---|
| | df | Plant-soil feedback | | Root | | |
| | | Shoot | | | | |
| | | | | | | |
| Species | 3 | 4.222 | 0.239 | | 5.358 | 0.147 |
| Nutrient | 1 | 0.122 | 0.727 | | 0.083 | 0.774 |
| Species x nutrient | 3 | 32.388 | <0.001 | | 4.882 | 0.181 |

Significant effects ($p < 0.05$) presented in bold.
Mycorrhizal fungi addition increased positive and decreased negative feedback

Plant-soil feedback of *Leontodon* and *Plantago* was, in addition to nutrient addition, also determined with addition of mycorrhizal spores at the seedlings. Relative colonisation of *Leontodon* roots by mycorrhizae was only significantly affected by nutrient status of the soil and not by mycorrhizal spore addition and soil history (Table S1; Fig. 5). *Leontodon* roots had a higher relative colonisation in high nutrient soils than in low nutrient soils (Fig. S5). Relative mycorrhizal fungi colonisation of *Plantago* roots, on the other hand, was significantly affected by both soil history and mycorrhizal spore addition (Table S1). On low nutrient soils, relative colonisation did not significantly differ between soil history and mycorrhizae treatments. On high nutrient soils, however, relative colonisation was higher with

Table 2 ANOVA results of nutrient and mycorrhizae (AMF) addition and their interaction on plant-soil feedback based on shoot and root biomass of *Leontodon* and *Plantago*

|             | Plant-soil feedback - *Leontodon* | Plant-soil feedback - *Plantago* |
|-------------|-----------------------------------|----------------------------------|
|             | Shoot | Root | Shoot | Root |
| df          | $\chi^2$ | $p$ | $\chi^2$ | $p$ | $\chi^2$ | $p$ | $\chi^2$ | $p$ |
| nutrient    | 1     | 11.358 | $<0.001$ | 9.737 | 0.002 |
| AMF         | 1     | 0.836 | 0.361 | 10.544 | 0.001 |
| nutrient x AMF | 1     | 0.792 | 0.373 | 1.686 | 0.194 | 8.207 | 0.004 | 3.504 | 0.061 |

Significant effects ($p < 0.05$) presented in bold
addition of mycorrhizal spores on both own and mix conditioned soil (Fig. S5).

Above- and belowground biomass of *Leontodon* and *Plantago* were only slightly affected by mycorrhizal spore addition (Fig. S6). Despite this and the small effects of mycorrhizal spore addition on visible mycorrhizal root colonisation, mycorrhizal spore addition did have significant effects on plant-soil feedback of the root system of *Leontodon* and interacted significantly with nutrient availability for plant-soil feedback based on the shoot of *Plantago* (Table 2; Fig. 3). *Leontodon* plant-soil feedback based on root biomass was positively affected by mycorrhizal spore addition and became significantly more positive on low nutrient soil, and the negative plant-soil feedback on high nutrient soil was significantly neutralised (Table 2; Fig. 3b). Plant-soil feedback based on shoot biomass of *Leontodon*, on the other hand, was not significantly affected by mycorrhizal spore addition (Table 2; Fig. 3a).

Discussion

Plant-soil feedback and nutrient availability

We found that plant-soil feedback based on shoot biomass, whether positive or negative, was eliminated with addition of nutrients. Species with neutral plant-soil feedback were not affected by nutrient addition. This is consistent with the general change in plant-soil feedback with nutrient addition that Petermann et al. (2008)

![Fig. 3](image-url) Plant-soil feedback based on (a) shoot and (b) root biomass of *Leontodon* and *Plantago* in low and high nutrient conditions, and with (blue) and without (yellow) addition of mycorrhizae spores (AMF) at seedling planting (n = 4–6). Different letters indicate significant differences (p < 0.05). Plant-soil feedback values without mycorrhizae spore addition are the same as in Fig. 2.
observed throughout 24 grassland species and the hypothesis of van der Heijden et al. (2008) that significance of microbial communities is highest at low nutrient availability. Our experimental approach, which quantifies the degree of microbial host specialisation (own versus a mixture of conditioned soil) (Lekberg et al. 2018), suggests that host specialisation was lost with nutrient addition. The neutralisation of the negative plant-soil feedback found for Plantago suggests that even though pathogens are often found to thrive in high nutrient soils (Paungfoo-Lonhienne et al. 2015; Whitaker et al. 2015; Wei et al. 2018), expected negative effects on plant growth (Rúa and Umbanhowar 2015; Revillini et al. 2016; van der Putten et al. 2016; Lekberg et al. 2018) were overcome. This possibly occurred due to increased overall fitness of the plant due to release from nutrient stress, and, consequently, an increase in tolerance to species-specific soil-borne pathogens. Simultaneously, competition between plants and soil micro-organisms may be less strong due to high nutrient availability (Schimel and Bennett 2004; Liu et al. 2016; ‘t Zandt et al. 2018), further decreasing negative plant-soil feedback effects.

For Leontodon, plant-soil feedback based on shoot biomass was positive under low nutrient conditions. Its elimination with nutrient addition may have occurred due to loss of mutualistic associations, i.e., positive associations were less frequently established or were unable to increase plant biomass any further than already occurred due to increased nutrient availability. Such decreased mutualistic associations with increasing nutrient availability have been shown to occur for mycorrhizal (Egerton-Warburton and Allen 2000; Corkidi et al. 2002; Treseder and Allen 2002; Treseder 2004; Avolio et al. 2014; Leff et al. 2015), Rhizobium (Vargas et al. 2000; Valladares et al. 2002) and plant growth-promoting rhizobacteria mutualisms (Shaharoona et al. 2008; Adesemoye et al. 2009).

Our results also show that nutrient availability effects of plant-soil feedback based on root system biomass do not match those of shoot biomass. In high nutrient soil, Leontodon showed an unexpected negative plant-soil feedback based on root biomass compared to no plant-soil feedback on low nutrient soil. A loss in significance of mutualistic associations together with a potentially flourishing pathogenic community (Paungfoo-Lonhienne et al. 2015; Whitaker et al. 2015; Wei et al. 2018; Veach et al. 2018) may have resulted in pathogenetic effects to prevail, and thus a negative plant-soil feedback on high nutrient soil. These negative effects were, however, not strong enough to show in the aboveground biomass, but may indicate that positive plant-soil feedback species experience small negative effects on own, conditioned soil with nutrient increase. Future research needs to establish whether this holds for positive feedback species in general and how this relation changes along a nutrient gradient.

Plant-soil feedback, mycorrhizal fungi interactions and nutrient availability

Mycorrhizal spore addition to the seedlings generally decreased negative and increased positive plant-soil feedback. This is consistent with the positive relation between plant-soil feedback and root mycorrhizal fungi colonisation on own conditioned soil that Cortois et al. (2016) found across 48 grassland species, and other studies experimentally stimulating mycorrhizal colonisation (Anacker et al. 2014; Garcia-Parisi and Omacini 2017). Our results show that these positive mycorrhizal effects interacted with soil nutrient availability. Negative feedback based on shoot biomass of Plantago lessened with mycorrhizal spore addition, but only on low nutrient soil. Also the increase in positive feedback of Leontodon with mycorrhizal spore addition only occurred on low nutrient soil. This is consistent with the notion that positive associations may become superfluous in high nutrient soil. Yet, for Leontodon, mycorrhizal fungi inoculation on high nutrient soil did affect plant-soil feedback based on root system biomass, and negative feedback became neutral. This together with the expectation that nutrient uptake was hardly limiting on high nutrient soil, suggests that the positive effect of mycorrhizal spore addition may result from protection against species-specific pathogens occurring on own soil (Sikes 2010), rather than an increased nutrient uptake via mycorrhizal networks.

Mycorrhizal fungi addition did not necessarily increase visible root colonisation, despite observed positive effects on plant-soil feedback. This is likely because root colonisation of mycorrhizal fungi is not the sole determinant of the functionality of the mutualistic interaction. Instead of the expected increase in mycorrhizal root colonisation, mycorrhizal spore addition may have resulted in more soil hyphae,
higher nutrient mobilisation capacity of the fungus and/or more nutrient exchange structures (Munkvold et al. 2004; Smith et al. 2004; Jansa et al. 2005; Feddermann et al. 2010). This may arise from an earlier developing plant-mycorrhizae association and result in the observed positive plant-soil feedback effects. Early effects of mycorrhizal fungi on plant-soil feedback direction and strength thus likely encompasses a much larger and complex pathway that is not captured in root colonisation determinants.

Soil nutrient increase and species co-existence theory

Microbial mediated plant-soil feedback is considered to play critical roles in plant community assembly via strong, negative effects on abundant species, resulting in reduced growth and subsequent local replacement (e.g. Bever et al. 2012). Such hypothesized continuous, small-scale species replacements may be affected by soil nutrient availability increase. If the neutralisation of plant-soil feedback with nutrient addition we observed occurs for all species in a diverse community, then continuous, small-scale species replacements may no longer take place in high nutrient environments. In this case, the best competitor will win and mono-dominate the system. Consequentially, high plant diversity can only be maintained in low nutrient environments, and will be lost with nutrient increase due to loss of microbial mediated plant-soil feedback. This is in line with observed loss of plant, bacterial and fungal diversity in the soil when soil nutrient availability increased (e.g., Clark and Tilman 2008; Smith et al. 2008; Coolon et al. 2013; Tedersoo et al. 2014). Future work should test whether root microbial communities between plant species in low nutrient soil are more distinct than in high nutrient soil, whether nutrient addition results in mono-dominance of strong competitor plant species, and whether this is linked to a loss of negative plant-soil feedback.

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Data availability The data used in these analyses are archived in the DANS-EASY data repository: https://doi.org/10.17026/dans-xas-6hts

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