

Nitrogen partitioning between plant species and soil microbes in alpine heath

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ABSTRACT

Nitrogen is rapidly taken up by plants and microbes, but questions remain as to which forms are preferred. Using *in situ* stable isotope labelling (¹³C and ¹⁵N), we show that co-existing plant species of alpine heath mainly take up ammonium and nitrate, passing ¹⁵N from root to shoot over time, leading to accumulated nitrogen in the shoots (over 10-fold increase compared with roots), with more complex organic nitrogen forms such as amino acids taken up to a lesser extent. Conversely, soil microbes preferred amino acids, potentially as a side-effect of satisfying their carbon requirements to build cellular structures. We show that competition for nitrogen can be alleviated by differing growth rates in plants and varying microbial preference of nitrogen forms.

Despite the prevailing idea that vascular plants prefer inorganic nitrogen (N), there is mounting evidence that they take up a range of N molecules, including amino acids (Näsholm et al., 1998; Ma et al., 2021; Liu et al., 2025). Past work shows that fast-growing (often dominant) plant species generally prefer inorganic N (Weigelt et al., 2005; Harrison et al., 2007), given that more complex organic forms may be energetically expensive to take up (Raab et al., 1999; McKane et al., 2002; Bardgett et al., 2003). Evidence also points to slower growing dwarf shrubs showing a stronger preference for more complex molecular forms, particularly in ecosystems at higher latitudes, likely due to their mycorrhizal associations, which are stronger at lower temperatures (Näsholm et al., 1998; Hu et al., 2024). Soil microbes likely also use more complex N forms than plants due to their capacity to produce enzymes capable of degrading complex compounds. Here, the presence of N in the molecule could be incidental. Microbial requirements for carbon mean the N is taken up as well, and often excreted as ammonium through mineralisation-immobilisation turnover (Barracough, 1997; Geisseler et al., 2009; Wilkinson et al., 2015). Therefore, inorganic N forms such as ammonium and nitrate may only be taken up by soil microbes when substrate C:N ratios are high (Hill and Jones, 2019). There is some evidence that microbes least prefer nitrate, with other N forms

taken up first (Barracough, 1997). However, in arctic and alpine ecosystems, low temperatures limit microbial activity in the soil, resulting in accumulation of organic matter and relatively high concentrations of dissolved organic N (Bardgett et al., 2007; Persson and Näsholm, 2001). Further, while microorganisms are highly competitive for N in the short-term, plants may be more competitive in the longer-term, but this is likely to be confounded by the enzymatic breakdown of complex amino acids into inorganic N (Harrison et al., 2007). Therefore, niche partitioning of N based on chemical form and temporal dynamics may facilitate plant-microbe coexistence (McKane et al., 2002), but substantial knowledge gaps remain as to how such processes operate *in situ*.

Here, we exposed co-existing alpine heath plant species, covering a range of growth rates and functional groups (dwarf-shrubs, graminoids and a forb), to mixtures of ¹⁵N-labelled inorganic and dual labelled (¹³C and ¹⁵N) amino acids of varying molecular weights (MW, Table 1). We hypothesised that 1) N uptake is partitioned between plants and microbes, where microbes take up amino acids but can break these down and make N available to plants over time; 2) soil microbes take up more carbon-rich, complex N forms than inorganic forms; and 3) plants generally prefer inorganic N forms over amino acids, but faster growing, more dominant species take up higher amounts of inorganic N, while

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Table 1
Chemical properties of the six N forms.

	Formula	C:N	Molecular weight
Ammonium	NH ₄	^a	18
Nitrate	NO ₃	^a	62
Glycine	C ₂ H ₅ NO ₂	2	75
Asparagine	C ₄ H ₈ N ₂ O ₃	2	132
Aspartic acid	C ₄ H ₇ NO ₄	4	133
Glutamic acid	C ₅ H ₉ NO ₄	5	147

^a Does not contain C.

slower growing dwarf shrubs may show a preference for amino acids.

The study was conducted in alpine heath on Glas Maol, Scotland (56°53'N, 3°22'W; 1000 m elevation; January/July average minimum/maximum temperatures: 5/-1 °C, 5/10 °C, respectively) in a plant community composed of, in increasing order of average dominance: *Galium saxatile* (forb), *Vaccinium vitis-idaea*, *V. myrtillus* (dwarf shrubs), *Carex bigelowii* (sedge) and *Festuca vivipara* (grass). In early spring, seventy plots (20 × 20 cm) were established across five blocks (14 plots per block) in a randomized block design (Fig. S1). Each plot contained all the aforementioned vascular plant species, which accounted for ~99% of total aboveground plant biomass within each plot. We created six solutions of 80 mg N L⁻¹ made up of a mixture of six equally weighted N forms (inorganic: ammonium [molecular weight 18], nitrate [62]; amino acids: glycine [75], asparagine [132], aspartic acid [133], glutamic acid [147]). The two inorganic N forms were labelled with 98% ¹⁵N, whereas the four amino acids were dual labelled with 98% ¹⁵N and ¹³C. A seventh, control solution was made using only unlabelled compounds. Direct uptake of the amino acids by plants would be indicated by plant tissue enrichment of both ¹³C and ¹⁵N (Näsholm et al., 1998). In June, one of the labelled solutions was added to two randomly selected

plots in each block. Each plot received 75 ml of labelled solution, via 25 × 3 ml injections in a 5 × 5 grid pattern across each plot, adding a total of 6 mg N to each plot. Using a soil bulk density of 0.35 g cm⁻³ (Ayres et al., 2006) and a plot size of 20 × 20 × 5 cm, this additional N is equivalent to 1.4 μg g⁻¹ soil. A preliminary analysis of soil N pools revealed 5.2, 8.0 and 12.0 μg g⁻¹ soil of nitrate, ammonium and dissolved organic N, respectively. Therefore, the amount of added N was unlikely to markedly influence plant or microbial growth (Ravn et al., 2017).

To assess the temporal fate of ¹⁵N and ¹³C, we collected turfs from the field experiment 2 and 45 days after labelling. At both sampling dates, seven turfs were taken from each block (one turf per labelled compound and the control). A 5 cm deep turf was removed from the central 15 × 15 cm of each plot and all vascular plant aboveground biomass was sorted by species within 24 h of turf collection, dried (60 °C, 48 h) and weighed to determine aboveground plant biomass. The soil was sieved (4 mm mesh) to collect the roots, and roots were dried (60 °C, 48 h) to determine community root biomass. Subsamples of shoot material from the five target species, as well as community root material, were ground for stable isotope analysis (Carlo-erba EA-DLT IRMS). The ¹⁵N and ¹³C in plant tissues were calculated according to standard methods, and correlations between ¹³C and ¹⁵N in each plant were assessed using Pearson's correlation coefficient to determine whether the molecule was taken up intact (Wanek and Arndt, 2002). Soil microbial biomass C and N were measured using fumigation-extraction (Brookes et al., 1985; Vance et al., 1987), and microbial biomass ¹⁵N concentrations were determined by diffusion of extracted N onto acidified paper discs (MacKown et al., 1987). Microbial uptake of ¹⁵N was calculated by dividing the excess ¹⁵N by the microbial biomass N. We calculated rate-based metrics for N uptake into plants by dividing the total excess ¹⁵N measured in each pool by the number of days from injection. ¹⁵N budgets were calculated by multiplying the concentration of excess ¹⁵N

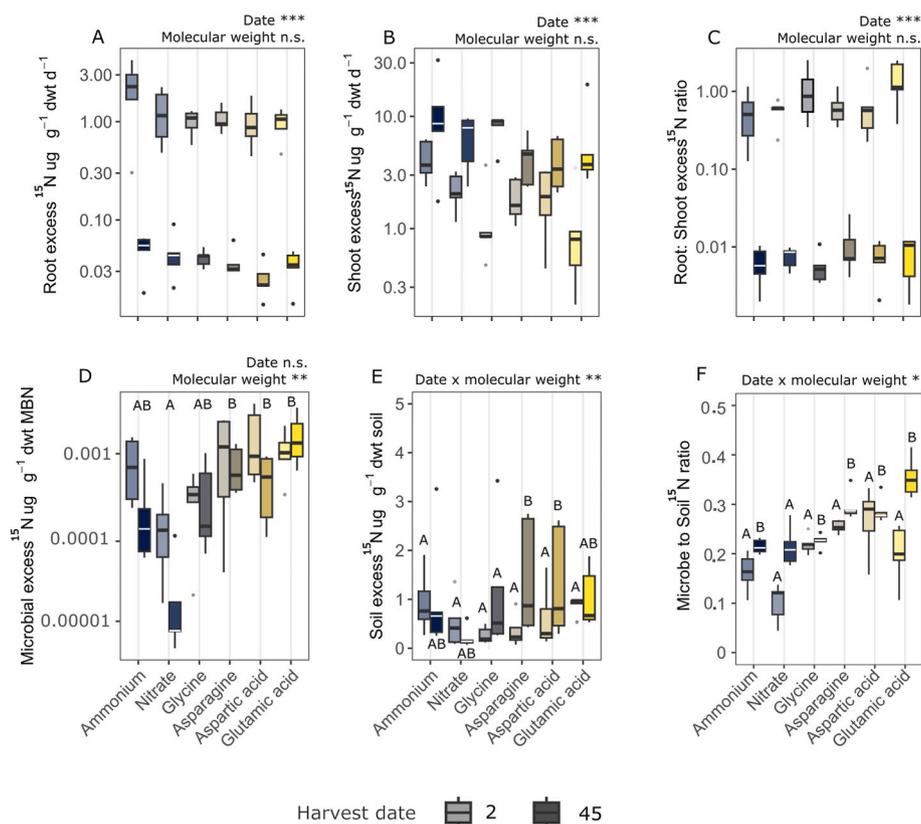


Fig. 1. Rate of uptake of different molecules with excess labelled ¹⁵N across harvest dates for A) root uptake over time (Harvest date: $F_{1,44} = 384.02$, $p < 0.001$), B) shoot uptake over time (Harvest date: $F_{1,44} = 234.26$, $p < 0.001$; Molecular weight: $F_{5,44} = 6.83$, $p < 0.001$), C) Root to shoot ¹⁵N ratio (Harvest date: $F_{1,44} = 58.95$, $p < 0.001$), D) microbial biomass ¹⁵N content (Molecular weight: $F_{5,44} = 4.34$, $p = 0.003$), E) bulk soil ¹⁵N content (Harvest date x Molecular weight: $F_{5,44} = 3.78$, $p = 0.006$), F) Microbial:Soil ¹⁵N (Harvest date x Molecular weight: $F_{5,44} = 3.39$, $p = 0.011$). Significance stars refer to: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

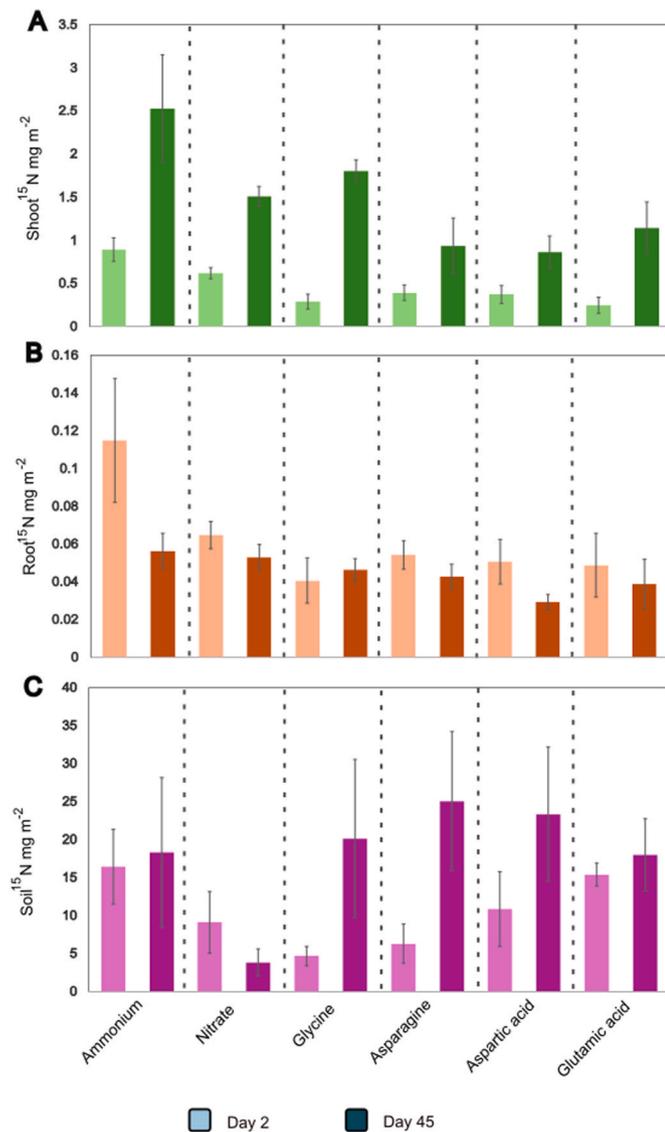


Fig. 2. Nitrogen budgets of the realised amounts of excess ^{15}N in shoots, roots and soil on days 2 and 45. The initial amount of ^{15}N added was 150 mg m^{-2} . Budgets were calculated by multiplying total biomass of each fraction, per turf, by the concentration of excess ^{15}N and scaling by m^2 . Soil mass is calculated to 5 cm deep at a bulk density of 0.35.

by the total turf level biomass of shoots, roots and microbes, or by the total weight of soil and scaling by area. Standard statistical analyses were then carried out (Text S1).

^{13}C uptake by plants was minimal, with some treatments losing ^{13}C relative to the background (range: $5 - 11\ \mu\text{g g}^{-1}$ shoot dry weight per day). ^{15}N uptake by roots slowed between days 2 and 45, as ^{15}N accumulated in the shoots (Fig. 1).

Rate of uptake of excess ^{15}N into the roots was significantly higher on Day 2 than 45 (Fig. 1A), but, while labelled amino acids appeared taken up at lower rates than labelled ammonium and nitrate, this was not statistically significant. The ^{15}N budget for roots shows that the total amount of labelled ammonium and nitrate decreased between days 2 and 45, and labelled glycine and asparagine increased (Fig. 2). Shoots accumulated excess ^{15}N at more rapid rates over the 45 days compared with the initial two days (Fig. 1B). In shoots there were also no effects of molecular weight, although again a slight downward trend with increasing molecular weight. The ^{15}N budget confirmed the increase in ^{15}N between days 2 and 45 in the shoots for all molecular forms (Fig. 2). Root to shoot ratios indicated greater (weight-corrected) retention by

roots relative to shoots at day 2 than day 45 ($F_{1,44} = 88.22$, $p < 0.001$, Fig. 1C). This suggests that roots were pushing ^{15}N rapidly through the plant regardless of initial labelled molecular type.

The correlation between ^{13}C and ^{15}N in shoot material could indicate direct uptake of the entire amino acid. However, we did not find strong evidence to suggest that large molecules were taken up intact by plants, and were more likely to have been mineralised by microbes and subsequently taken up by plants in inorganic forms (negative correlation; Table S1).

Excess ^{15}N in the microbial biomass was not affected by harvest date, potentially indicating low turnover of the ^{15}N through microbes (Fig. 1D). Retention of labelled nitrate by microbes was significantly lower than the three largest amino acids (Fig. 1D and 2). Further, there was a decline in nitrate, and to some extent ammonium, in microbial biomass between days 2 and 45. Nitrate must be reduced to ammonium through nitrate reductase before it can be used by microbes (Pan et al., 2023). Not all microbial taxa synthesise microbial reductase and for those that do, the energetic cost is high (Hiis et al., 2024). Therefore, nitrate uptake is often associated with low concentrations of ammonium in the soil (González et al., 2006). While ammonium is thought to be the preferred N form for microbial uptake, this tends to be the case when carbon rich molecules are abundant and was not the case in our study, as evidenced by the comparative rate of uptake of amino acids (Farrell et al., 2014). Additionally, ammonium can only be used by microbes if there are many carbon sources available to maintain cellular processes (Wilkinson et al., 2014). Taken together, it seems that in our study carbohydrates were low in the soil, meaning that nitrate and ammonium could not be effectively used by the microbes and were therefore discarded by day 45.

Soil excess ^{15}N showed an interaction between date and molecular weight. The data in Fig. 1E shows that labelled N from both asparagine and aspartic acid accumulated in the soil on day 45 and were significantly higher than at day 2 and also accumulated in higher amounts than all the other molecular forms (Fig. 1E). The ^{15}N budget shows that much labelled N was returned to the soil by day 45, which could indicate the microbial community had broken down the amino acids and retained the C, discarding excess N. The ratio of ^{15}N uptake in microbes to soils also showed an interaction between date and molecular weight (Fig. 1F). On day 2 all ratios were the same between molecules, while on day 45 labelled ammonium, glycine, asparagine, aspartic acid and glutamic acid were significantly higher in microbes than the soil compared with nitrate. This trend indicated a clear preference by microbes for amino acids. Soil microbes are often C-limited, rather than N-limited due to generally low concentrations of available organic matter in soil and its physical and chemical protection (Lehmann and Kleber, 2015; Soong et al., 2020) and microbes synthesise enzymes capable of breaking down more complex C- and N-containing compounds (Daly et al., 2021). Our finding that microbes take up amino acids and retain them in their biomass, while nitrate uptake is extremely low, could be indicative of preference for amino acids. The fate of labelled nitrate that was lost from microbes between days 2 and 45, also seen in the ^{15}N budget, is not explained by uptake into plant tissues or released to bulk soil (Fig. 2). Therefore, it is possible nitrate was denitrified to nitrogen oxide or leached to deeper soil layers so the N is not retained within microbial biomass. Nitrate leaching is one of the first signs of saturation of N in an ecosystem and is commonly observed in heathland (Phoenix et al., 2012). The ^{15}N budget supports the likelihood of leaching and possibly denitrification, as it shows very low levels of labelled nitrate in all compartments on both days 2 and 45 (Fig. 2).

There are likely several factors contributing to the low recovery of all ^{15}N forms observed in our study (150 mg m^{-2} added compared with a maximum of 30 mg m^{-2} total recovered on days 2 and 45), which is lower than reported in other labelling studies using similar approaches (Bardgett et al., 2003; Harrison et al., 2007). Potential reasons for this low recovery of ^{15}N include the leaching or N to deeper soil layers and uptake by bryophytes, which wasn't accounted for in this study. Indeed,

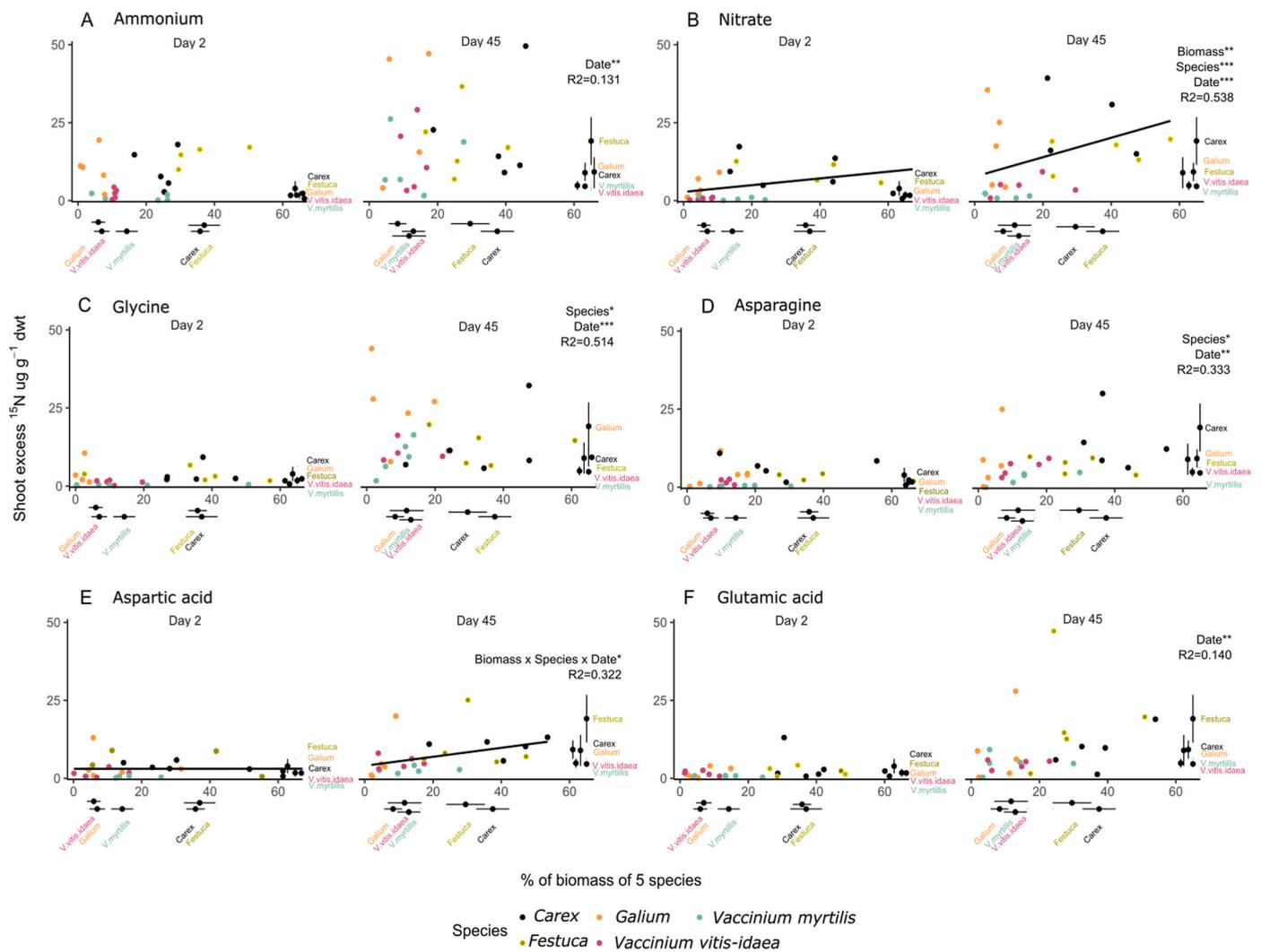


Fig. 3. Relationships between proportion of aboveground biomass of the five plant species and labelled ^{15}N uptake in the shoot biomass of each species on Day 2 and Day 45, presented by molecular weight of labelled N species. The black lines show the overall community trend where significant. The overarching model indicated highly significant interactions between harvest date and molecular weight: $F_{5,171} = 3.43$, $p = 0.006$; harvest date and biomass share: $F_{1,171} = 7.81$, $p = 0.006$; and molecular weight and species: $F_{20,171} = 2.08$, $p = 0.006$. See Table S3 for a simplified overview of ^{15}N uptake patterns and molecular weight-specific model output. The black points outside the graphs show the mean of the biomass share against the y axis and nitrogen uptake against the x axis for each species, with error bars representing standard error of the mean.

studies on fate of N indicate that N is commonly lost to deeper soil layers and that bryophytes can directly uptake ^{15}N from soil in these arctic (Ayres et al., 2006) and other heathlands (Ray, 2007; Phoenix et al., 2012). Albeit over longer timescales, other studies have also found comparably low retention of ^{15}N in the top 5 cm of soil (~20%) in the year after labelling (Andresen et al., 2023).

We observed a strong interactive effect between harvest date and the species' proportional share of community biomass on the concentration of shoot ^{15}N , partially supporting our third hypothesis that fast growing, dominant plant species would take up N more rapidly than less dominant species (Fig. 3). On day 2, there was generally a neutral or weakly positive relationship between excess ^{15}N and increasing proportion of community biomass, while on day 45 the relationship tended towards positive or neutral except for glycine, which was slightly negative (Fig. 3). We did not find significant relationships between individual plant species abundance and shoot uptake of ^{15}N , likely due to the small sample size. The two most dominant species in every turf were the sedge *Carex bigelowii* and the grass *Festuca vivipara*. In the case of ammonium, nitrate, glycine, aspartic acid and glutamic acid, where one species (e.g. *Carex*) was more dominant, the other, less dominant species (e.g. *Festuca*) took up more N, potentially indicating competition between the

two species. Further, the forb *Galium saxatile* had very low relative abundance, often the lowest in the turf, but took up disproportionately high amounts of N for all forms, suggesting it is a highly acquisitive species. Our expectation in the third hypothesis was that the two *Vaccinium* species, as dwarf shrubs, would take up amino acids preferentially, but we found little evidence of that. Hill and Jones (2019) discussed the difficulty in identifying uptake of intact amino acids and suggested that a problem in definitively tracking ^{13}C could arise when background ^{13}C is high, which is potentially the case in alpine systems where decomposition is slow.

The partitioning of N forms by plants and soil microbes is a topic that has received relatively little attention, despite its potential importance for efficient N use in N limited ecosystems, such as alpine heath as studied here. Our findings indicate that over time ^{15}N from both inorganic forms and amino acids are retained in microbial and plant tissues. Fast-growing plants generally take up simpler forms of N and vascular plants and microbes have different preferences for chemical forms of N. We also found that plants that make up the highest proportion of community biomass tend to take up more ^{15}N , but such biomass-N uptake relationships can differ at the community versus species level (Shen et al., 2024).

Here we showed that in alpine heath, partitioning of N based on chemical form occurs between plants and microbes. While we did not find evidence to determine whether amino acids were taken up by plants and microbes intact, we detected a clear preference for amino acids by microbes likely because of low availability of carbon. This is likely to arise through chemical or physical protection of soil organic matter from decomposition. Further, while higher plant biomass at a species level was associated with greater uptake for nitrate and aspartic acid, we showed that the two most dominant plant species did not follow the biomass-ratio hypothesis, which could indicate high plant-plant competition for N in these alpine heaths. The cold climate and low nutrient availability of these heathlands results in complex uptake patterns for plants and microbes as colimitation of nutrients, and competition for resources, result in constraints in effectively using nutrients when they become available.

CRedit authorship contribution statement

Ellen L. Fry: Writing – review & editing, Writing – original draft, Visualization, Formal analysis. **Jonathan R. De Long:** Writing – original draft, Visualization, Investigation, Formal analysis. **Edward Ayres:** Methodology, Investigation, Data curation, Conceptualization. **René van der Wal:** Writing – review & editing, Writing – original draft, Resources, Project administration, Methodology, Investigation, Funding acquisition, Formal analysis, Conceptualization. **Richard D. Bardgett:** Writing – review & editing, Writing – original draft, Supervision, Resources, Project administration, Methodology, Investigation, Funding acquisition, Data curation, Conceptualization.

Declaration of competing interest

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: Richard Bardgett, René Van der Wal reports financial support was provided by Lancaster University. If there are other authors, they declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.soilbio.2026.110127>.

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