

Plant and microbial N acquisition under elevated atmospheric CO₂ in two mesocosm experiments with annual grasses

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Abstract

The impact of elevated CO₂ on terrestrial ecosystem C balance, both in sign or magnitude, is not clear because the resulting alterations in C input, plant nutrient demand and water use efficiency often have contrasting impacts on microbial decomposition processes. One major source of uncertainty stems from the impact of elevated CO₂ on N availability to plants and microbes. We examined the effects of atmospheric CO₂ enrichment (ambient + 370 μmol mol⁻¹) on plant and microbial N acquisition in two different mesocosm experiments, using model plant species of annual grasses of *Avena barbata* and *A. fatua*, respectively. The *A. barbata* experiment was conducted in a N-poor sandy loam and the *A. fatua* experiment was on a N-rich clayey loam. Plant–microbial N partitioning was examined through determining the distribution of a ¹⁵N tracer. In the *A. barbata* experiment, ¹⁵N tracer was introduced to a field labeling experiment in the previous year so that ¹⁵N predominantly existed in nonextractable soil pools. In the *A. fatua* experiment, ¹⁵N was introduced in a mineral solution [(¹⁵NH₄)₂SO₄ solution] during the growing season of *A. fatua*. Results of both N budget and ¹⁵N tracer analyses indicated that elevated CO₂ increased plant N acquisition from the soil. In the *A. barbata* experiment, elevated CO₂ increased plant biomass N by ca. 10% but there was no corresponding decrease in soil extractable N, suggesting that plants might have obtained N from the nonextractable organic N pool because of enhanced microbial activity. In the *A. fatua* experiment, however, the CO₂-led increase in plant biomass N was statistically equal to the reduction in soil extractable N. Although atmospheric CO₂ enrichment enhanced microbial biomass C under *A. barbata* or microbial activity (respiration) under *A. fatua*, it had no significant effect on microbial biomass N in either experiment. Elevated CO₂ increased the colonization of *A. fatua* roots by arbuscular mycorrhizal fungi, which coincided with the enhancement of plant competitiveness for soluble soil N. Together, these results suggest that elevated CO₂ may tighten N cycling through facilitating plant N acquisition. However, it is unknown to what degree results from these short-term microcosm experiments can be extrapolated to field conditions. Long-term studies in less-disturbed soils are needed to determine whether CO₂-enhancement of plant N acquisition can significantly relieve N limitation over plant growth in an elevated CO₂ environment.

Keywords: *Avena barbata*, *Avena fatua*, elevated CO₂, microbial biomass C and N, ¹⁵N availability, N tracer, plant–microbial N partitioning

Received 17 May 2004; revised version received 20 August 2004 and accepted 11 October 2004

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Introduction

The concentration of atmospheric CO₂ has increased by nearly 100 ppm since preindustrial times (Keeling *et al.*, 1989) and is predicted to double within the next century

largely because of the increasing combustion of fossil fuel and land use changes (IPCC 2001). CO₂ is an important greenhouse gas contributing 55% of all the greenhouse gas forcing in the 1980s, and is estimated to have an even greater relative importance in the 1990s through to the 21st century (Lashof & Ahuja, 1990). Terrestrial ecosystems contain nearly three times as much C (ca. 2060 Gt C) as that currently present in the atmosphere (ca. 735 Gt C) (Houghton & Woodwell, 1989) and have the potential to be either a significant C sink or C source under future CO₂ scenarios (Raich & Potter, 1995; Hungate *et al.*, 2003). Hence there is increasing interest in the possibility of modulating atmospheric CO₂ concentration via enhanced below ground carbon storage in managed terrestrial systems. The feasibility of this approach will depend to a large degree on our ability to successfully manage the 'plant C-fixation–microbial decomposition loop'. There has now been a substantial amount of work on elevated CO₂ and plant response. Over the short-term, elevated CO₂ often stimulates photosynthesis, enhances plant nutrient demand and nutrient use efficiency, and improves plant water use efficiency (Coleman & Bazzaz, 1992; Jackson *et al.*, 1994). This can affect ecosystem C balance by increasing net primary production (NPP) (Melillo *et al.*, 1993) and modifying decomposition processes (Ball, 1997). However, the long-term impacts on ecosystem C balance are not clear because those CO₂-induced alterations may have interactive, often contrasting, effects on soil microbial processes, particularly on soil N immobilization and mineralization (Richter *et al.*, 2003).

The primary productivity of most temperate terrestrial ecosystems is N-limited (Vitousek & Howarth, 1991); hence a major source of uncertainty stems from the effect of CO₂ elevation on plant and microbial N acquisition and subsequent N cycling, which may significantly impact further ecosystem responses to CO₂ enrichment (Daepf *et al.*, 2000; Schlesinger & Lichter, 2001; Gill *et al.*, 2002; Luo *et al.*, 2004). On the one hand, elevated CO₂ often increases NPP and litter C/N or lignin/N ratios, favoring C or N accumulation in plant biomass and soil organic matter (Rouhier *et al.*, 1994; Oren *et al.*, 2001), and slowing nutrient cycling (Cotrufo & Ineson, 1996; Hu *et al.*, 2001). On the other hand, elevated CO₂ may enhance root exudation and C availability for microbes, stimulating microbial activity and N mineralization, and increasing the proportion of C and N undergoing rapid cycling (Zak *et al.*, 1993; Hungate *et al.*, 1997; Paterson *et al.*, 1997; Pendall *et al.*, 2004). Also, CO₂-led increases in plant C:N ratios are likely species-dependent and less significant in naturally senesced litters of tree species (Norby *et al.*, 2001). Whether CO₂-enhancement of plant photosynthesis

and subsequent C accumulation can be sustained will largely depend on the potential for microbes to obtain and/or release limited nutrients to balance the needs for continuous plant growth.

Studies on the effects of elevated CO₂ on microbial processes have generally emphasized the importance of enhanced C-input to microbial biomass and activity (Diaz *et al.*, 1993; Zak *et al.*, 1993; Paterson *et al.*, 1997; Hu *et al.*, 1999), assuming that microbes in soil are C-limited (Smith & Paul, 1990). However, both plants and microbes are commonly N-limited in terrestrial ecosystems (Kaye & Hart, 1997) and litter decomposition can be inhibited by plant competition for N (Wang & Bakken, 1997). CO₂ enrichment in the atmosphere alters resource availability (particularly the relative availability of C and N) for microbes, potentially altering the microbial community composition and functions (Montealegre *et al.*, 2002). The vast majority of N in soil exists in the form of organic N with low C:N ratios ranging from 10–12:1 in agricultural soils to 20–25:1 in forest soils (Fog, 1988). Therefore, one important question is whether or not the generally enhanced C availability for microbes under elevated CO₂ stimulates microbial utilization of this old organic N (Zak *et al.*, 1993; Cardon *et al.*, 2001; Hu *et al.*, 2001; Richter *et al.*, 2003).

Little attention has been directed toward the effects of elevated CO₂ on plant–microbial competition for N and the subsequent effects on microbial N acquisition from old organic matter. Decomposition experiments, largely conducted in soils without the presence of actively growing plants, indicate that compared with ambient controls, the decomposition rate of litter produced under elevated CO₂ is slower over the short-term (Cotrufo & Ineson, 1996; Ball, 1997). On one hand, elevated CO₂ might promote microbial N utilization and subsequent immobilization because of increased C supplies to microbes (Diaz *et al.*, 1993). On the other hand, it may promote plant N acquisition through enhancing the growth of fine roots and mycorrhizae (Rillig *et al.*, 1999; Hu *et al.*, 2001). In addition, results from other experiments suggest that enhanced microbial activities induced by elevated CO₂ may increase N mineralization (Zak *et al.*, 1993; Hungate *et al.*, 1997). However, clear information addressing the effects of elevated CO₂ on microbial N acquisition in the presence of active plants is very limited. This knowledge gap severely limits our ability to predict ecosystem C balances under the future conditions of elevated atmospheric CO₂ (Niklaus *et al.*, 2001).

In this paper, we present data obtained from two microcosm experiments using ¹⁵N as a tracer to examine the effects of elevated CO₂ on plant–microbial N partitioning and plant N acquisition. Our objectives were (1) to examine whether elevated CO₂ increases

plant and microbial N acquisition, and (2) to determine where the increased N in biomass originates if plant and microbes enhance their N uptake.

Materials and methods

Two growth chamber experiments were conducted to determine the effects of elevated CO₂ on plant-microbial N partitioning. In the first experiment, mineral ¹⁵N was introduced in the previous season (1997) in a field labeling experiment, and ¹⁵N was distributed among various soil pools (Hu *et al.*, 2001). In the second experiment, ¹⁵N was introduced during the growing season as soluble mineral N. In addition, soil texture and fertility were different between the two soils used, with experiment 1 on a nutrient-poor sandy loam and experiment 2 on a relatively N-rich clay loam. The two model plant species used, annual graminoid *Avena barbata* and *A. fatua*, are common in California grasslands.

Experiment 1

Experimental setup. The first experiment was conducted in regular growth chambers at the University of California at Berkeley, CA. The soil was collected from an isotope (¹⁵N)-labeled experiment at the Jasper Ridge Biological Preserve, Palo Alto, CA, USA (Hu *et al.*, 2001). The field site was an annual grassland dominated by *A. barbata*. The ¹⁵N tracer was introduced into the soil on 1 March, 1997 by injecting (¹⁵NH₄)₂SO₄ solution (4.0 mg ¹⁵N kg⁻¹ soil) into a depth of about 10 cm. Soils (15 cm depth) were collected when plants senesced on 23 April, 1997 (Hu *et al.*, 2001). The sandstone grassland soil contained 1.68% total organic C and 0.17% organic N as quantified after plant materials (both shoots and roots) were removed after the ¹⁵N labeling experiment. The soil was sieved (2 mm mesh) and residues on the sieve were removed, leading to very low C availability for soil microbes. Because of the ¹⁵N-labeling treatment in the previous season, the soil had a higher ¹⁵N content equivalent to an addition of 2.66 μg ¹⁵N g⁻¹ soil. The ¹⁵N largely existed in nonextractable pools (73%) and microbial biomass (21.7%), with a very small fraction in extractable mineral N (5.3%) (Hu *et al.*, 2001). The soil had been kept at 4 °C for about 16 months until being used for this experiment.

Annual graminoid *A. barbata* seeds were collected from the Jasper Ridge Biological Preserve, Palo Alto, CA. Seeds of *A. barbata*, a nonnative annual grass, were pregerminated at room temperature for 48 h and two healthy germinating seeds were planted into each pot microcosm (10 cm × 10 cm × 10 cm). Each microcosm

contained exactly 670 g soil (dry-soil equivalent). These microcosms were immediately transferred into growth chambers with a 12 h photoperiod, 25 °C day, 15 °C night, and 70% relative humidity. Two treatments, ambient and elevated CO₂ (ambient + 370 μmol mol⁻¹), were applied 1 week after seeding, with 10 replicates for each treatment. The microcosms were rotated biweekly between the two chambers to eliminate any potential chamber effects (Diaz *et al.*, 1993) and microcosms were rotated within each chamber. During the whole growing season, the same amount of water was added to each pot using a syringe as needed to ensure adequate water was available (as shown by no plant withering) and watering was carefully conducted to ensure no water leakage. A saucer was also placed under each microcosm pot to further ensure no leaking of water occurred.

Sample collection and soil, plant and microbial analyses. Microcosm pots were destructively harvested when seeds of *A. barbata* matured on 16 March, 1999. Plant shoots and roots were manually separated and the soil samples were sieved (2 mm mesh). Roots on the sieve were manually sorted, washed thoroughly and oven-dried (65 °C, 72 h).

Microbial respiration in root-free soils was measured as total CO₂ evolution from incubation at 22 °C for 7 days at constant moisture (15% w/w, ca. 75% field water holding capacity). Microbial biomass C and N were determined by fumigation-extraction (Vance *et al.*, 1987), using 0.5 M K₂SO₄ extraction after being shaken for 30 min. Soil extractable organic C in the K₂SO₄ extracts before and after the fumigation was quantified using a total C analyzer (Shimadzu TOC-5050A, Shimadzu Co., Kyoto, Japan). Soil extractable inorganic N (NH₄-N and NO₃-N) in the nonfumigated and fumigated soils was measured on a flow injection analyzer (Lachat Quickchem Systems, Milwaukee, WI, USA) after alkaline persulfate digestion (Cabrera & Beare, 1993). The total of extractable NH₄⁺ and NO₃⁻ in K₂SO₄ extracts of nonfumigated samples was defined as extractable N.

The C and N concentrations of soil, shoots and roots and their isotope ratios were determined by combustion on a gas chromatograph-mass spectrometer (GC-MS) (Europa Scientific LTD., Crewe, UK), using finely ground subsamples. The ¹⁵N ratio in the extractable pool of N was determined following diffusion and the microbial biomass ¹⁵N ratio was determined following Kjeldahl digestion and diffusion (at University of California at Berkeley), and the diffusion process was carried out by using filter paper disks in PTFE (Teflon) traps in plastic specimen cups (Stark & Hart, 1996). Sample δ¹⁵N (‰) values were converted to excess

nitrogen isotope (mg) and conversion of $\delta^{15}\text{N}$ (‰) to the absolute isotope ratio ($^{15}\text{N}/^{14}\text{N}$) of the sample was based on the atom ratio of atmospheric nitrogen. Sample ^{15}N content was then calculated from fractional abundance ($^{15}\text{N}/(^{15}\text{N} + ^{14}\text{N})$) and total N content of the sample. Nonextractable ^{15}N content (mostly organic ^{15}N) was calculated by subtracting microbial biomass ^{15}N and K_2SO_4 -extractable ^{15}N from total soil ^{15}N .

Experiment 2

The second experiment was conducted at the USDA Air-Quality CO_2 Facility, North Carolina State University, Raleigh, NC, USA. The facility consists of a $9\text{ m} \times 12\text{ m}$ bay and contains 20 continuous stirred tank reactor (CSTR) chambers for exposure of plants to CO_2 (Booker *et al.*, 2000). The chambers were built in a regular greenhouse and no extra light was applied. Each CSTR (1.2 m diameter \times 1.4 m tall) can be used to expose plants to CO_2 . Air sampling is accomplished by extracting a small quantity of gas exiting the CSTR and drawing it back through Teflon lines into the laboratory. A computer activates and deactivates solenoid valves (one per CSTR) to sample the air in each chamber for approximately 2 min and averages and collects CO_2 data for analysis.

Experimental setup. The soil was a fine-loam over clayey, mixed, mesic Ultic Haploxeralf (Sutherlin series) collected from an annual grassland at the University of California Hopland Research Station in Mendocino County, CA, USA ($39^\circ 00'\text{N}$ latitude, $123^\circ 4'\text{W}$ longitude). About 80 kg soil (sampled to 10 cm) was obtained, sieved (4 mm mesh), and well mixed before being shipped to North Carolina State University, Raleigh, NC, USA. The soil had been kept under 4°C for 18 months until being used for this experiment. Two CO_2 concentrations, ambient and elevated (ambient + $370\ \mu\text{mol mol}^{-1}$), were applied, with three replicates (i.e. three CSTR chambers) for each CO_2 treatment. In each CSTR chamber, six microcosm pots ($12\text{ cm} \times 12\text{ cm} \times 15\text{ cm}$) were randomly placed and their positions were frequently rotated. Each pot contained exactly 1294.36 g of soil (dry soil equivalent). Twenty-five pregerminated *A. fatua* seeds (48 h) were planted into each pot and then thinned 1 week later so that each pot had 20 healthy plants and the CO_2 treatments immediately started thereafter. During the whole growing season, watering was carefully conducted as described in Experiment 1.

Introduction of ^{15}N tracer. Exactly 4.0 mg ^{15}N ($^{15}\text{NH}_4)_2\text{SO}_4$, 99.7% atom ^{15}N) per kg of soil were injected into soils on 10 March, 2001 in 60.0 mL of H_2O solution,

introducing a total of 5.177 mg ^{15}N into each microcosm pot at six points to a 10 cm depth into the soil. Watering was carefully conducted to ensure no water leakage and associated ^{15}N leaching.

Collection and analyses of plant and soil samples. Microcosm pots were destructively harvested on 11 March, 24 April and 23 May (i.e. 1, 45 and 75 days after the ^{15}N introduction). These sampling dates corresponded approximately to the late tillering, flowering and maturing stages of *A. fatua*. The last sampling date physiologically corresponded to the harvest date for *A. barbata* in experiment 1. Microbial respiration in root-free soils was measured as total CO_2 evolution from incubation at 22°C for 7 days at constant moisture (18% w/w, equivalent to 70–75% water holding capacity) (Hu & van Bruggen, 1997). Microbial biomass C and N were determined by fumigation–extraction as previously described. Plant shoot, root and soil subsamples were oven-dried (65°C , 72 h) and finely ground using a ball mill. The C and N concentrations of soil, shoots, and roots were quantified on a Perkin-Elmer 2400 CHNS/O elemental analyzer (Norwalk, CT, USA) and their isotope ratios were determined by combustion on a ThermoFinnigan DELTAPlus continuous flow isotope ratio mass spectrometer (CF-IRMS) (ThermoFinnigan, Bremen, Germany), using finely ground subsamples (measured at the Soil Analytical Lab., North Carolina State University, Raleigh, NC, USA). Microbial biomass ^{15}N ratio was determined following persulfate digestion and diffusion (Stark & Hart, 1996) and ^{15}N in the nondigested samples were taken as the ^{15}N of the extractable inorganic N. Sample $\delta^{15}\text{N}$ (‰) and sample ^{15}N content were calculated as described above.

Mycorrhizal infection of plant roots. The percentage of root length colonized by arbuscular mycorrhizal (AM) fungi was measured on roots stained in trypan blue (Phillips & Hayman, 1970) using the gridline-intersect method (Giovannetti & Mosse, 1980). Briefly, roots (about 1 cm long) were cleared in KOH, acidified in HCl, and then stained with trypan blue solution. The stained roots were spread on a Petri dish with gridlines, and examined on a dissecting microscope at $\times 40$ magnification. The intersections between roots and gridlines were counted. The percentage of the colonized roots was then calculated.

Data analyses. Plant biomass (root and shoot) and biomass N was converted to a base unit of per m^2 of soil. One-way variance analysis (ANOVA) was used to detect the effects of elevated CO_2 on plant shoot and root biomass, biomass N and ^{15}N , microbial biomass C, microbial biomass N, microbial respiration, and soil

extractable inorganic N ($P \leq 0.05$). For all statistical analyses, the SPSS V.10.0 (SPSS Inc., Chicago, IL, USA) software package was used.

Results

Shoot and root biomass and biomass N

Elevated CO₂ in the atmosphere significantly increased plant biomass in both *A. barbata* and *A. fatua* experiments (Fig. 1). In the first experiment, elevated CO₂ significantly increased *A. barbata* shoot biomass by 36.5% from 235 to 320 g m⁻² (Fig. 1a) and shoot biomass N by 25.1% from 1.3 to 1.6 g N m⁻² (Fig. 1b). However, root biomass (299 and 338 g m⁻², respectively) and biomass N (1.4 and 1.3 g m⁻², respectively) were not significantly different between the control and elevated CO₂. Still, CO₂ enrichment significantly increased total plant biomass N by 9.8%. In addition, elevated CO₂ significantly increased biomass C:N ratios from 73 to 80 in shoots and 85 to 100 in roots. By the final harvest, root biomass constituted ca. 50% of the total plant biomass in *A. barbata*.

In the second experiment, CO₂ enrichment significantly increased *A. fatua* shoot biomass by 11%, 53% and 42% at the March, April and May sampling dates, respectively (Fig. 1c). However it significantly increased total N in shoots only in the April and May samples (10.5% and 12.9% for shoots, respectively) (Fig. 1d). The final shoot biomass N was 7.1 and 8.1 g N m⁻² soil for the control and elevated CO₂ treatments, respectively. Also, CO₂ enrichment significantly increased root biomass (77% and 60%, respectively) and biomass N (23% and 36%, respectively) in the April and May samples (data not shown). The root biomass N at the final harvest was 1.7 and 2.4 g N m⁻² soil for the ambient control and elevated CO₂ treatment, respectively. Biomass C:N ratio significantly increased from 52 to 59 in roots and 55 to 63 in shoots from the control to elevated CO₂. By the final harvest, root biomass constituted less than 25% of the total plant biomass in *A. fatua*. In addition, total *A. fatua* biomass did not increase between the April and May sampling, but root biomass was slightly reduced. By the final harvest, total plant biomass N of *A. fatua* was 13.6 g m⁻² soil under elevated CO₂, 17.8% higher than the ambient control (11.3 g m⁻² soil).

K₂SO₄ extractable N in soil. Elevated CO₂ did not significantly reduce soil extractable N in the *A. barbata* experiment. Although the mean value of the extractable N was ca. 16% lower in the elevated CO₂ treatment (0.71 mg N kg⁻¹ soil) than in the ambient control (0.85 mg N kg⁻¹ soil), this CO₂-induced reduction was

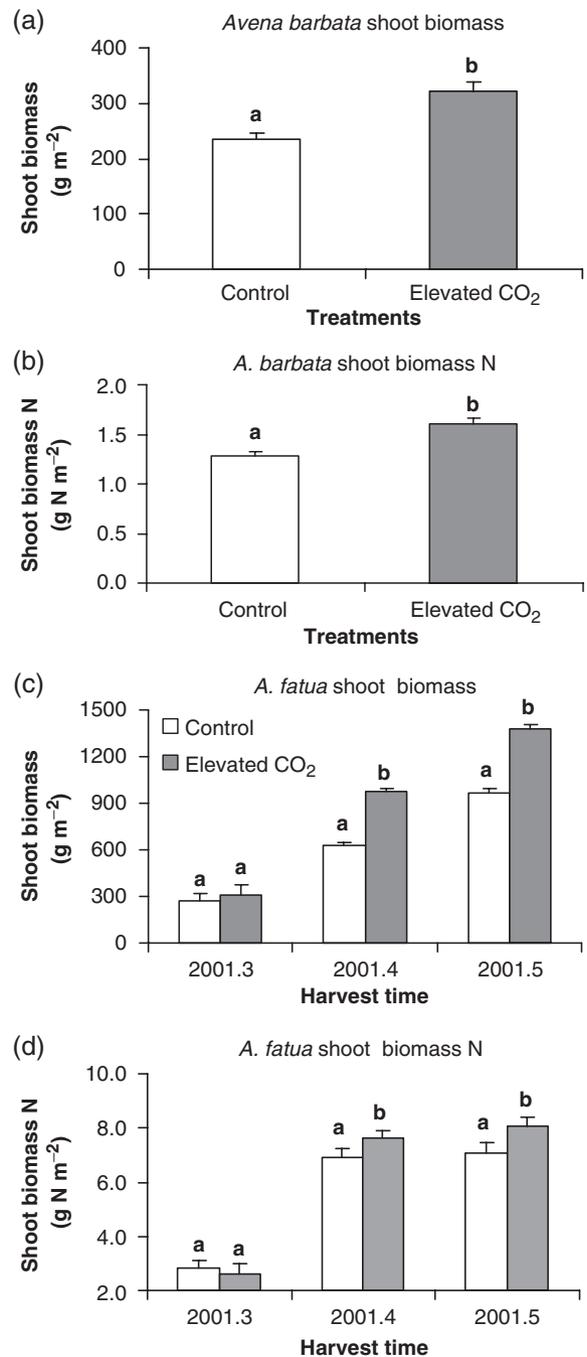


Fig. 1 The effects of CO₂ enrichment on plant biomass and biomass N of *Avena barbata* (a, b) and *A. fatua* (c, d). The unit is g m⁻² for plant biomass and g N m⁻² for plant biomass N. Bars depict treatment means and standard errors (SEM). Bars with the same letter at the sampling date are not significantly different at $P \leq 0.05$.

not statistically significant (Fig. 2a) and only accounted for ca. 5% of the increased biomass N. At the early stage of the *A. fatua* experiment when plants were small,

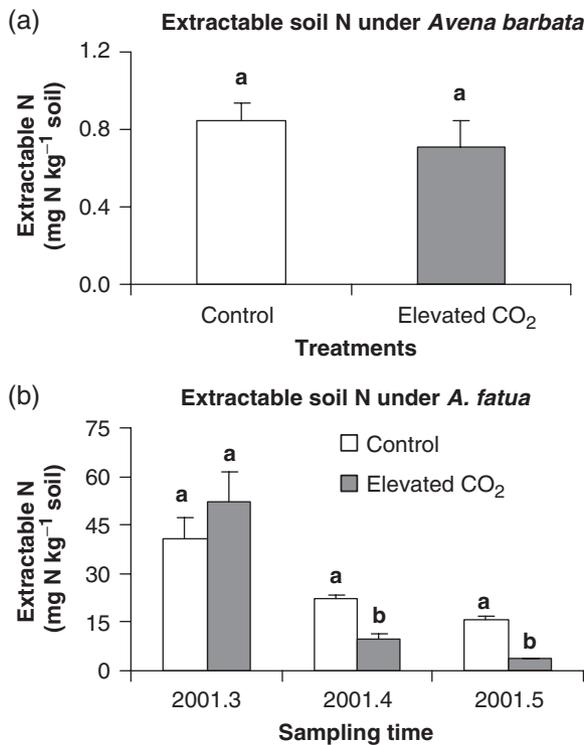


Fig. 2 Elevated CO₂ effects on soil extractable N under *Avena barbata* (a) and *A. fatua* (b). Bars depict treatment means and standard errors (SEM). Bars with the same letter at the sampling date are not significantly different at $P \leq 0.05$.

elevated CO₂ did not affect soil extractable N, as shown by samples collected on 11 March. But as the growing season advanced, CO₂-induced decreases in extractable N became significant. By the final harvest time, total extractable N was averaged at 3.84 mg N kg⁻¹ soil in the elevated CO₂ pots, in comparison with 15.99 mg N kg⁻¹ soil in the ambient control (Fig. 2b). The reduction in extractable N in soil (averaged at 12.15 mg N kg⁻¹ soil) under elevated CO₂ was approximately equal to the increase in total plant biomass N (averaged at 12.24 mg N kg⁻¹ soil).

Soil microbial biomass carbon and nitrogen. Microbial biomass C (fumigation flush) was significantly higher under elevated CO₂ in the *A. barbata* experiment (Fig. 3a), but microbial biomass N (fumigation flush) showed no difference between the ambient control (18.3 mg N kg⁻¹ soil) and the elevated CO₂ (19.3 mg N kg⁻¹ soil). In the *A. fatua* experiment, CO₂ enrichment did not increase microbial biomass C (Fig. 3b) but enhanced microbial respiration in the late growing stage (Fig. 3c). Microbial biomass N showed no significant difference between the treatments and among the sampling dates, averaging at 30.3 mg N kg⁻¹ soil in the ambient control and 28.1 mg N kg⁻¹ soil under elevated CO₂.

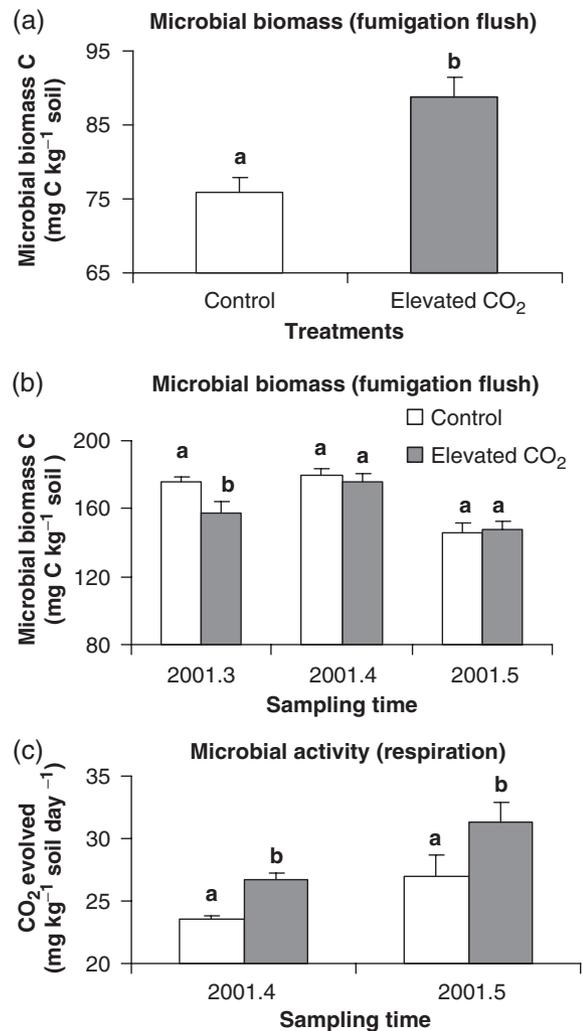


Fig. 3 Elevated CO₂ effects on soil microbial biomass C and microbial activity under *Avena barbata* (a) and *A. fatua* (b, c). Bars depict treatment means and standard errors (SEM). Bars with the same letter at the sampling date are not significantly different at $P \leq 0.05$.

¹⁵N distribution among plant, soil and soil microbes. Elevated CO₂ altered ¹⁵N distribution among plant shoots and roots, soil and soil microbes (Figs 4 and 5). Total ¹⁵N in *A. barbata* shoots and roots was not significantly impacted by CO₂ elevation (data not shown). However, by the final harvest, percentage recovery of introduced ¹⁵N in *A. fatua* materials increased from 2.6% to 4.6% in roots, and from 18.2% to 26.0% in shoots under elevated CO₂ (Figs 4a and b). Elevated CO₂ in the atmosphere reduced the ¹⁵N concentration of *A. barbata* shoots and roots because of dilution effects of increased biomass, but increased both shoot and root ¹⁵N concentrations in *A. fatua* (Table 1). ¹⁵N in the extractable N pool was not different between the ambient control and elevated CO₂ in the *A. barbata*

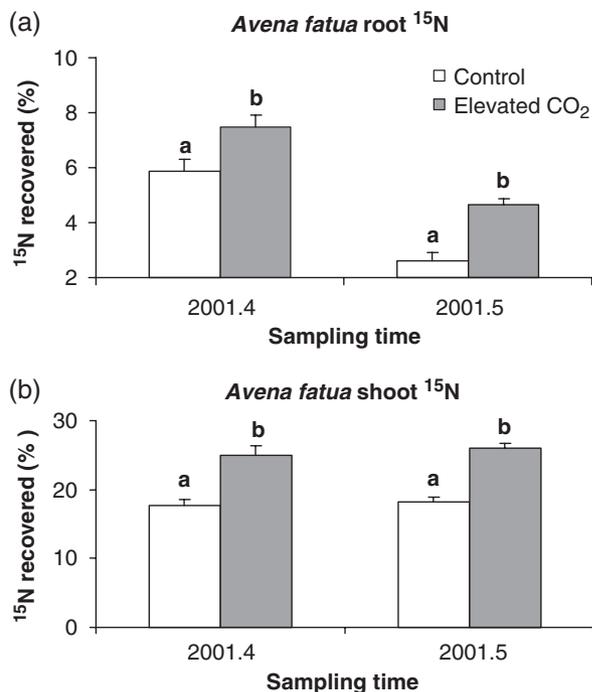


Fig. 4 Elevated CO₂ effects on plant ¹⁵N acquisition of *Avena fatua*: shoot ¹⁵N (a) and root ¹⁵N (b). Bars depict treatment means and standard errors (SEM). Bars with different letters at the sampling date are significantly different at $P \leq 0.05$.

experiment (data not shown). However, in the *A. fatua* experiment, total ¹⁵N in soil (% recovery) was significantly reduced with elevated CO₂ by the late growing season to 45.5%, compared with 49.4% in the ambient control (Fig. 5a). The reduction was highly significant in the extractable N pool (Fig. 5b). By the final harvest time, about 22% of introduced ¹⁵N remained in the extractable soil N pool in the control, but this number reduced to 6.0% under elevated CO₂. Total recovery of ¹⁵N (i.e. the sum of ¹⁵N in soil and plant shoots and roots) was significantly higher in the elevated CO₂ (76% of total ¹⁵N introduced) than in the control (70.2% of total ¹⁵N introduced). Total ¹⁵N in the microbial biomass N (ca. 1% of the total ¹⁵N introduced) and microbial biomass ¹⁵N:¹⁴N ratios were not impacted by elevated CO₂ in either experiment (data not shown).

Mycorrhizal colonization of A. fatua roots. Mycorrhizal colonization of *A. fatua* roots was significantly higher under elevated CO₂ than in the ambient control (Fig. 6). CO₂-led increases in root mycorrhizal colonization were about 71% in the April samples and 48% in the May samples. For the samples collected at the final harvest, the magnitude of increases in mycorrhizal infection (48%) was close to that in aboveground plant

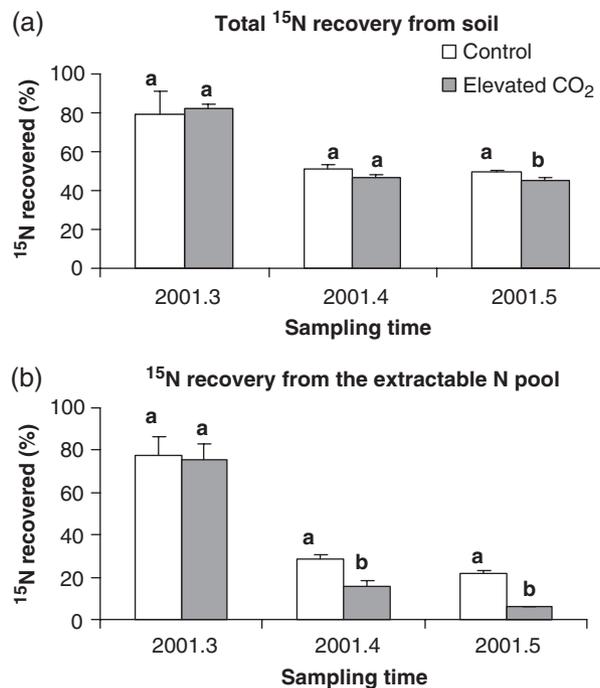


Fig. 5 Effects of CO₂ enrichment on soil ¹⁵N distribution under *Avena fatua*: (a) total ¹⁵N in soil, and (b) ¹⁵N recovered from the extractable soil N pool. Bars depict treatment means and standard errors (SEM). Bars with the same letter at the sampling date are not significantly different at $P \leq 0.05$.

biomass (42%, Fig. 1). However, there was no correlation between these two parameters ($R^2 = 0.03$; $P > 0.05$). Over time, root mycorrhizal colonization significantly increased from April (10.8% in the ambient control and 18.5% in the elevated CO₂) to May (21.0% and 31.1%, respectively).

Discussion

Results obtained from our experiments indicated that elevated CO₂ led to increased plant N uptake in both highly N-limiting and relatively N-rich soils (Fig. 1). However, microbial biomass N did not increase under elevated CO₂, in spite of increases in microbial biomass C under *A. barbata* (Fig. 3a) or microbial activity under *A. fatua* (Fig. 3c), suggesting differential effects of CO₂ enrichment on plant and microbial N acquisition.

The data from the ¹⁵N tracer provided some direct evidence showing that elevated CO₂ increased N transfer from the soil to plants (Figs 1, 2, 4 and 5). These results were consistent with our previous observations in a grassland dominated by *A. barbata* (Hu *et al.*, 2001), suggesting that elevated CO₂ may favor plants over microbes for N acquisition. Similar results have recently been reported in several long-term

Table 1 Elevated CO₂ effects on ¹⁵N concentrations (¹⁵N/(¹⁵N + ¹⁴N) × 100) of plant and soil samples[†]

CO ₂ treatment	Sampling date					
	1		2		3	
	Ambient	Elevated	Ambient	Elevated	Ambient	Elevated
<i>Experiment 1 (Avena barbata)</i>						
Shoots	ND	ND	ND	ND	1.24	1.08*
Roots					1.12	0.97*
Soil					0.49	0.49
<i>Experiment 2 (A. fatua)</i>						
Shoots	0.49	0.54	1.78	2.12*	1.77	2.09*
Roots	0.62	0.65	1.29	1.29	1.20	1.32*
Soil	0.54	0.54	0.48	0.47	0.48	0.46*

*Significant difference ($P \leq 0.05$) between the control and elevated CO₂ at its corresponding sampling date.

[†]Sampling dates 1, 2 and 3 approximately correspond to tillering, flowering and maturing stages of *A. barbata* (March 16, 1999 only) and *A. fatua* (March 11, April 23 and May 24, 2001, respectively).

ND means no data because plants and soils were not sampled.

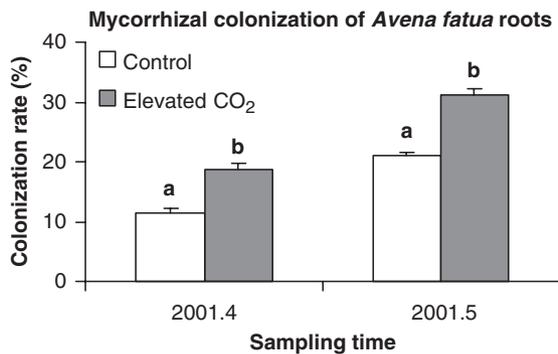


Fig. 6 Elevated CO₂ effects on mycorrhizal colonization of roots in *Avena fatua*. Bars depict treatment means and standard errors (SEM). Bars with different letters at the sampling date are significantly different at $P \leq 0.05$.

field experiments. For example, Richter *et al.* (2003) reported that in *Lolium perenne* and *Trifolium repens* swards, microbial N immobilization was not strongly affected by 7-years of exposure to elevated CO₂. Niklaus *et al.* (2003) also observed no changes in microbial biomass N in a nutrient-poor grassland after being exposed to elevated CO₂ for six growing seasons. Together, these results suggest that enhanced microbial immobilization may not be a major mechanism constraining plant response to elevated CO₂, as initially proposed (Diaz *et al.*, 1993). The implications for this alteration of N partitioning between plants and microbes are not exactly known. However, an understanding of the mechanisms that govern CO₂ enhancement of plant N acquisition may help us predict the long-term consequences. Experimental evidence demonstrating CO₂-led increases in root biomass and C

inputs is plentiful (Treseder & Allen, 2000; BassiriRad *et al.*, 2001; Niklaus *et al.*, 2003; Treseder *et al.*, 2003; Pendall *et al.*, 2004), similar to the CO₂-led increases in root biomass and mycorrhizal infection in our current experiments. This evidence indicates that plants under elevated CO₂ may be able to adjust their C allocation to exploit N (and possibly other nutrient) resources in the soil (i.e. compensatory adjustments to increase acquisition capacity for minerals, BassiriRad *et al.*, 2001; Pendall *et al.*, 2004). However, what is unknown is the potential of CO₂-enhancement of plant N uptake and where the enhanced N, if any, originates (Richter *et al.*, 2003).

In many temperate ecosystems, the soil contains a total organic N that is at least two magnitudes larger than the annual needs of plants. Still, plant growth in these ecosystems is most likely N-limited (Vitousek & Howarth, 1991) because soil microbes do not release the N contained in plant residues and other older organic matter to permit plants uptake. Since elevated CO₂ often stimulates microbial activity through increasing C inputs to soil, it has been proposed that CO₂-enhancement of microbial activities would increase N mineralization, thereby sustaining N supply for plants (Zak *et al.*, 1993; Hungate *et al.*, 1997). A gradual conversion of this organic N into plant biomass with high C:N would sustain plant N requirements under elevated CO₂ over the long term, without the need for new N inputs. Results from our *A. barbata* experiment seemed to support this hypothesis as the CO₂-led increase in plant biomass N cannot be balanced unless microbes released some N from other organic N. CO₂ enhancement of plant biomass N but not total ¹⁵N in *A. barbata* suggest that microbes might have released some N

from older organic pools. However, the data obtained from our *A. fatua* experiment was inconsistent with this hypothesis. There, enhanced N uptake under elevated CO₂ predominantly originated from the extractable N pool as increases in plant biomass N were accompanied by comparable decreases in extractable N. Atmospheric CO₂ enrichment only facilitated the transfer of extractable N to plants as evidenced by increased ¹⁵N concentrations in *A. fatua* tissues. These results are consistent with those obtained from open-top chambers in the field by Cardon *et al.* (2001), who showed that CO₂-enhanced microbial activity as measured by respiration may largely stem from preferential consumption of newly fixed organic C, rather than N-enriched old organic materials. Similarly, Richter *et al.* (2003) examined gross N fluxes in a grassland soil exposed to elevated atmospheric CO₂ for 7 years and found that organic N turnover and microbial N mineralization were not affected. The difference in the results between our two experiments may be related to the initial nutrient status, particularly the N availability. Elevated CO₂ may prompt plants to acquire N in the extractable pool first before microbes access old organic N (if any). The magnitude of microbial conversion of older organic N under elevated CO₂ may be critically important in understanding plant and microbial response to progressive N limitation under elevated CO₂ (Hungate *et al.*, 2003; Luo *et al.*, 2004).

One interesting finding in our experiment is that CO₂-enhancement of mycorrhizal colonization of roots (Fig. 6) coincided with a marked decrease in soil extractable N in the *A. fatua* experiment (Fig. 2). Stimulation of mycorrhizae by elevated CO₂ has been documented in various experiments (Rillig *et al.*, 1999; Treseder & Allen, 2000; Treseder *et al.*, 2003). However, increased plant N uptake has not been attributed directly to the enhanced mycorrhizae. Instead, enhancement of plant N uptake has previously been attributed to higher soil moisture and extended plant growing periods under elevated CO₂ (Jackson *et al.*, 1994), which is highly possible in some water-limiting systems (Hu *et al.*, 2001; Billings *et al.*, 2002). However, frequent watering was applied to avoid water limitation for the plants in our experiments. Also, plant roots essentially occupied the whole volume of the soil in our confined systems and mycorrhizal contribution for N uptake was expected to be minimal if their role is mainly extension of root surface. It is unknown why plant roots did not effectively uptake extractable N in the ambient control, although by the middle of the growing season plant leaves became yellowish in both treatments (similar to a N-limiting symptom). Rapid depletion of extractable N under elevated CO₂ suggests that mycorrhizal hyphae may still be able to reach N pools that plant roots are

unable to. Alternatively, CO₂-enhancement of mycorrhizae may have stimulated plant uptake of other limiting nutrients such as P (Grunzweig & Korner, 2003), leading to a corresponding increase of plant N utilization. CO₂-stimulation of mycorrhizae and its resulting effects on N and P availability may have some long-term implications in mediating plant and ecosystem response to rising CO₂ in the atmosphere. In grasslands where AM fungi dominate, enhancement of mycorrhizae is unlikely to directly stimulate organic matter decomposition as AM fungi generally lack saprotrophic capacity (Allen, 1991; Read & Perez-Moreno, 2003). However, increased C inputs associated with enhanced mycorrhizal hyphae may indirectly stimulate residue decomposition by increasing microbial activity (Hodge *et al.*, 2001; Pendall *et al.*, 2004; Tu *et al.*, unpublished data). Whether the resulting N release from organic matter can partially relieve N constraints on long-term plant and ecosystem responses to rising atmospheric CO₂ levels deserves more attention.

In summary, results obtained from our microcosm experiments showed that elevated CO₂ increased plant N acquisition without significantly impacting microbial biomass N. In the first experiment with a N-poor soil, the CO₂-enhanced N obtained by plants largely originated from the nonextractable pools. In the second experiment with a N-rich soil, it predominantly stemmed from the extractable N pool. However, the mechanisms that underlie the enhancement of plant N uptake are unclear and it is unknown whether results obtained in these short-term microcosm experiments can be extrapolated to the field. Potential effects of CO₂-enhancement of mycorrhizae on plant N and P acquisition warrant further investigation.

Acknowledgements

We thank H. L. Zhong, Donald Herman and Howard Sanford for ¹⁵N measurements, Valerie Eviner, Christine Johnson and Karen Parker for experiment execution and I-Long Wu for sample preparation. Thanks were also extended to Jeff Barton for his technical help in maintaining the CSTR facility and to the USDA Air Quality group (Raleigh, NC) for access to the CSTR facility. We highly appreciate four anonymous reviewers for their comments on an earlier version of the manuscript. The experiments were partially supported by grants from the Energy Foundation of the University of California, the National Science Foundation (DEB 9627368 and 01686), and USDA-NRI.

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