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Regenerating temperate forest mesocosms in elevated CO₂: belowground growth and nitrogen cycling

Received: 10 January 1997 / Accepted: 25 July 1997

Abstract The response of temperate forest ecosystems to elevated atmospheric CO₂ concentrations is important because these ecosystems represent a significant component of the global carbon cycle. Two important but not well understood processes which elevated CO₂ may substantially alter in these systems are regeneration and nitrogen cycling. If elevated CO₂ leads to changes in species composition in regenerating forest communities then the structure and function of these ecosystems may be affected. In most temperate forests, nitrogen appears to be a limiting nutrient. If elevated CO₂ leads to reductions in nitrogen cycling through increased sequestration of nitrogen in plant biomass or reductions in mineralization rates, long-term forest productivity may be constrained. To study these processes, we established mesocosms of regenerating forest communities in controlled environments maintained at either ambient (375 ppm) or elevated (700 ppm) CO₂ concentrations. Mesocosms were constructed from intact monoliths of organic forest soil. We maintained these mesocosms for 2 years without any external inputs of nitrogen and allowed the plants naturally present as seeds and rhizomes to regenerate. We used ¹⁵N pool dilution techniques to quantify nitrogen fluxes within the mesocosms at the end of the 2 years. Elevated atmospheric CO₂ concentration significantly affected a number of plant and soil processes in the experimental regenerating forest mesocosms. These changes included increases in total plant biomass production, plant C/N ratios, ectomycorrhizal colonization of tree fine roots, changes in tree fine root

architecture, and decreases in plant NH₄⁺ uptake rates, gross NH₄⁺ mineralization rates, and gross NH₄⁺ consumption rates. In addition, there was a shift in the relative biomass contribution of the two dominant regenerating tree species; the proportion of total biomass contributed by white birch (*Betula papyrifera*) decreased and the proportion of total biomass contributed by yellow birch (*B. alleghaniensis*) increased. However, elevated CO₂ had no significant effect on the total amount of nitrogen in plant and soil microbial biomass. In this study we observed a suite of effects due to elevated CO₂, some of which could lead to increases in potential long term growth responses to elevated CO₂, other to decreases. The reduced plant NH₄⁺ uptake rates we observed are consistent with reduced NH₄⁺ availability due to reduced gross mineralization rates. Reduced NH₄⁺ mineralization rates are consistent with the increases in C/N ratios we observed for leaf and fine root material. Together, these data suggest the positive increases in plant root architectural parameters and mycorrhizal colonization may not be as important as the potential negative effects of reduced nitrogen availability through decreased decomposition rates in a future atmosphere with elevated CO₂.

Key words *Betula* · CO₂ · Mycorrhizal fungi · Nitrogen · Pool dilution

Introduction

The large and rapid rise in atmospheric CO₂ concentrations (Keeling and Whorf 1994) has the potential of directly altering earth's climate system (Schneider 1989) and net primary productivity (NPP) of terrestrial ecosystems (Melillo et al. 1996). The direct effects of elevated CO₂ on terrestrial ecosystems is important for two reasons. First, increased plant growth in elevated CO₂ may serve as an important negative feedback on the rate of rise of atmospheric CO₂ levels (Schimel 1995). Second, altered plant growth and physiology in elevated

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CO₂ atmosphere may lead to significant changes in the structure and functioning of managed and natural ecosystems (Bazzaz 1990; Bazzaz and Sombroek 1996; Koch and Mooney 1996).

Temperate forest ecosystems are likely to play a key role in mediating future climate change. Recent theoretical and empirical analyses have suggested that the terrestrial biosphere, especially temperate forest ecosystems, are a globally important sink for atmospheric CO₂ (Ciais et al. 1995; Goulden et al. 1996; Tans et al. 1990, 1995; Wofsy et al. 1993). Current estimates suggest that, on average, forest species respond to a doubling of atmospheric CO₂ concentrations with about a 30% increase in NPP, (Wullschleger et al. 1995).

In addition to potential effects on NPP, elevated CO₂ may lead to changes in the composition of forest ecosystems through the differential effect of elevated CO₂ on the growth of different species (Ceulemans and Mousseau 1994; Poorter 1993; Wullschleger et al. 1995). Changes in species composition in an elevated CO₂ environment are likely to be especially important in regenerating forest ecosystems, which themselves are likely to be common within future landscapes (Bazzaz et al. 1996). There are a number of factors which may increase the frequency of incidents of forest regeneration in the future. These include changes in temperature and precipitation which may lead to shifts in tree species ranges (Davis 1989; Davis and Zabinski 1992); altered climatic regimes which could enhance disturbance and/or forest dieback through the increased frequency of pest outbreaks and/or forest fires (Kurz et al. 1995); and an increased frequency of extreme weather events which could lead to increased disturbance (Bassow et al. 1994; Katz and Brown 1992).

Much of our current understanding of how different species respond to elevated CO₂ are derived from studies of individually grown plants. Unfortunately, individually grown plants appear to respond to elevated CO₂ quite differently than plants grown in more natural, interacting assemblages (Ackerly and Bazzaz 1995; Bazzaz and McConnaughay 1992; Wayne and Bazzaz 1995). Natural assemblages of plants, and their associated natural soil environments, are increasingly being used to study plant community responses to elevated CO₂ (e.g., Arp et al. 1993; Díaz et al. 1993; Field et al. 1996; Körner et al. 1996; Navas et al. 1995; Owensby et al. 1994). This approach allows us to study an integrated set of interactions (plant-plant and plant-soil) which are critical components of natural ecosystem function.

Our current understanding of the direct effects of elevated CO₂ on aboveground growth and physiology is better developed than for belowground growth and physiology (Rogers et al. 1994). However, belowground plant growth responses and interactions with the soil environment may play a critical role in maintaining or constraining long-term plant growth responses to elevated CO₂ (Curtis et al. 1994). These belowground feedbacks consist of indirect effects of elevated CO₂ on nutrient cycling in the soil and direct effects on plant

root architecture and mycorrhizal symbioses (Norby 1994; O'Neill 1994). Indirect effects of elevated CO₂ on nutrient cycling in the soil can be either positive or negative. Increases in plant C/N ratios can lead to reduced rates of mineralization in the soil (Berntson and Bazzaz 1997). Increases in carbon allocation belowground may either increase or decrease nutrient availability through changes in the soil microbial population (Díaz et al. 1993; Zak et al. 1993). And finally, changes in plant water use efficiency may increase soil water content which in turn enhances mineralization rates (Hungate et al. 1996). Changes in plant root architecture and mycorrhizal symbioses almost always lead to increased potential nutrient acquisition by plants and thus represent a mechanism whereby plants in elevated CO₂ atmospheres may be able to acquire more nutrients and thus maintain growth enhancement (Berntson 1994; Norby 1994; O'Neill 1994). To identify the relative importance of these multiple belowground feedbacks and mechanisms of sustained growth enhancement, it is important that they be simultaneously characterized within integrated experiments of terrestrial ecosystem responses to elevated CO₂ (Berntson and Bazzaz 1996a).

In this paper, we present a study of the impact of elevated CO₂ atmospheres on the first two years of regeneration in mesocosms consisting of intact monoliths of forest soil. Using this experimental system of a regenerating temperate forest, we test the following hypotheses regarding the effects of elevated CO₂:

1. Plant species composition is changed.
2. Belowground plant growth and mycorrhizal symbioses are enhanced.
3. Nitrogen cycling within the soil, including plant and microbial uptake of nitrogen, is changed.

Together, we assess the potential relative importance of these responses (especially 2 and 3) in relation to the availability and uptake of nitrogen by regenerating forest communities in elevated CO₂ atmospheres.

Materials and methods

Intact monoliths of forest floor soil (Oe + Oa) measuring 25 × 40 cm were collected at the Harvard Forest Long Term Ecological Research Site (Petersham, Mass.) in early May 1994. This site is located in the Transition Hardwood-White Pine-Hemlock forest region of New England (Westveld et al. 1956). Forest floor material was collected from several locations within an area of approximately 360 m² where the dominant canopy trees were (in order of decreasing percent basal area) *Quercus rubra*, *Acer rubrum*, *Pinus strobus*, *Betula alleghaniensis*, *B. papyrifera*, and *Prunus serotina* (Bassow 1995). We removed undecayed leaf litter from the monoliths.

Monoliths were carefully transported back to a controlled growth facility in Cambridge, Massachusetts, cut to size, and placed within plastic containers (0.325 × 0.25 × 0.25 m, 20 l) with drainage holes. The top of the monoliths was within a cm of the top of the tub. The bottom of the tubs contained a coarse silica sand which provided ample drainage and no additional nutrients. The total time between soil collection from the field and placement into the containers was less than 48 h. The average depth of the forest floor material was just over 7 cm and settled an insignificant amount (<0.5 cm) through the duration of the experiment.

The plants present within the seed and rhizome bank within the soil were allowed to regenerate without interference. The mesocosms were never fertilized, but they were watered every 1–2 days so the regenerating plants were never water stressed. All senesced leaves were allowed to fall to the soil within each tub.

A total of 24 mesocosms were distributed among six controlled-environment glasshouses, and placed under 30% shade cloth to simulate a diffuse canopy, but otherwise received natural lighting. Half of the glasshouses (3 units) were maintained with ambient atmospheric CO₂ (375 ppm) and the other with elevated CO₂ (700 ppm). Day/night temperatures were maintained at 26°/19°C with a 13-h day until the beginning of September. At this time day/night temperatures were slowly ramped down to simulate the onset of autumn and harden off the plants for winter. At the end of the first growing season (November 1994), when >90% of leaves had senesced and the plants were dormant, the tubs were moved to an outside garden for the remaining winter months. Tub was placed in trenches so that their soil surface was even with the external soil surface, and soil was filled in around the outside of each tub to equalize soil temperatures between the inside and outside of the tubs. The tubs were brought back into the glasshouses at the first sign of bud swelling (mid-March 1995).

¹⁵N-NH₄⁺ tracer application and measurement

In the second week of August 1995 (year 2), 120 ml of 2.5 mmol l⁻¹ 98% ¹⁵N-NH₄Cl was evenly distributed (vertically and horizontally) within each mesocosm by 120 1-ml injections using a specially built injection system (Berntson 1996). We selected this level of tracer addition because it represented approximately a 25% increase above ambient soil NH₄⁺ levels measured in the field (Aber et al. 1993). Soil samples were collected immediately prior to injection, and 15 min, 24 h, and 48 h following injection (4 sampling times). Four cores of soil (1 cm diameter, full depth of forest floor) were collected at randomly selected locations within each mesocosm and pooled together. ¹⁵N injections, soil processing, and plant harvesting were processed in three blocks (paired ambient + elevated CO₂ environments, four replicates in each) over a 6-day period.

Soil samples were extracted in 0.5 M K₂SO₄ (Davidson et al. 1989) with a soil/extract ratios (w/v) of 0.1. K₂SO₄ solutions were filtered on pre-combusted glass fiber filters. Samples were placed on a shaker for 24 h prior to filtration, and then frozen for storage. Concentrations of NH₄⁺ in the soil extracts were determined using a Westco Scientific auto analyzer (Westco Scientific, Danbury, Conn.) using a modified Berthelot reaction with phenol (Maynard and Kalra 1993).

The level of ¹⁵N enrichment (%¹⁵N atoms) for the NH₄⁺ recovered in the soil extracts taken prior to, after 15 min, 24 h, and 48 h after injection of the label was determined by diffusion to 0.625-cm-diameter pre-combusted glass fiber filters (Gelman AE). The filters contained 20 µl 2 M H₂SO₄ and were wrapped in polytetrafluoroethylene (Sørensen and Jensen 1991). To volatilize the NH₄⁺, 0.25 g MgO was added to the extracts (80 ml) in air-tight urine specimen containers. To ensure that the MgO was sufficiently mixed within the K₂SO₄ extracts, acid washed boiling chips were added and the specimen containers were placed on a gently moving shaker table for 10 days. Glass fiber filters were dried in a desiccator which contained a concentrated H₂SO₄ trap to remove contamination by atmospheric NH₄⁺. Blanks consisted of glass fiber filters with 20 µl 2 M H₂SO₄ which were placed in the desiccator with the samples. Determination of the total N and ¹⁵N content of the glass fiber filters was performed using a Europa Scientific ANCA-sl coupled to a model 20–20 Stable Isotope Analyzer (Europa Instruments, England).

Soil microbial biomass was determined by chloroform fumigation (Jenkinson and Powlson 1976) of soil samples followed by direct extraction in 0.5 M K₂SO₄ (Brookes et al. 1985; Davidson et al. 1989). This procedure was performed only on the soil samples collected 48 h after ¹⁵N injection. For each these samples, half of the soil was directly extracted in K₂SO₄, and the other half was first fumigated with chloroform. Total-N and ¹⁵N enrichment in the

K₂SO₄ solutions were determined before and after fumigation. Measurement of C, N and ¹⁵N content in the K₂SO₄ solutions was performed using the Europa ANCA-sl and 20–20 stable isotope analyzer. Dehydrated K₂SO₄ (to which no soil had been added) was used as a blank. To maximize the amount of solution which could be processed, the K₂SO₄ solutions were dehydrated by placing them in a drying oven for 24 h at 75°C. Samples of the dehydrated K₂SO₄ (75–150 mg) were then processed using the Europa ANCA-sl and stable isotope analyzer described above. To obtain maximum resolution when processing these samples an offset O₂ injection was used, providing resolution down to 0.5 µmol N (Owens and Rees 1989). The amount of ¹⁵N recovered in microbial biomass was corrected by assuming that natural ¹⁵N abundance of the microbial pool was equal to the natural ¹⁵N abundance of the soil NH₄⁺ pool.

Plant harvesting, mycorrhizal assessments and quantification of root architecture

After the 48-h incubation with the ¹⁵N-NH₄ tracer, each mesocosm was harvested and all the plant material carefully collected. At the start of harvest, all plant stems were cut approximately 1 cm above the soil surface and tags were attached to the remaining stems so that each root system could be identified to species. The roots of all the plants for each different species of plant encountered were carefully washed from the soil by placing the entire soil block in a tub of water, and then carefully disentangling and extracting the roots by hand. It was inevitable that some of the fine roots were lost, but care was taken to keep each root system as intact as possible. Entire root systems for each species were wrapped in a moist paper towel, placed in an air tight plastic bag (with all the air removed) and frozen prior to the morphological and mycorrhizal measurements. Above-ground plant material was oven-dried for 96 h at 70°C and weighed to the nearest mg.

To characterize the root morphology and ectomycorrhizal status of each species, we made detailed measurements on subsamples of the fine roots of each species from each mesocosm. For all of the tree species, fine roots were defined as roots <0.5 mm in diameter which showed no clear signs of secondary growth or lignification. Woody roots consisted of those with clear secondary growth and lignification. For the herbaceous species, no attempt was made to separate different fractions of the root systems or characterize ectomycorrhizae. A total of approximately 200 individual root tips (215 ± 11, mean ± SE) were taken from three or four randomly selected locations within each root system. For each subsample, the total number of root tips, and for the tree species, the total number of root tips with a mycorrhizal sheath, were counted under a dissecting microscope. Root tips which were broken or dead were not included in these counts. Once the root tips had been counted, the subsample was laid out in a petri dish with ~2mm of water to minimize overlap of individual roots and scanned into a computer as 8-bit gray scale images at 177.2 pixels cm⁻¹. From these images total root length and average diameter of the fine root subsample were calculated using the procedures and algorithms proposed by Pan and Bolton (1991). From all of these measurements made on the (fine) root subsamples the following parameters were derived: percent mycorrhizal root tips, root diameter, specific root length (SRL, m g⁻¹) and specific root number (SRN, number of root tips g⁻¹). Total fine root length and number of root tips for each species within a given mesocosm were derived by multiplying SRL and SRN by (fine) root mass.

After taking the fine root subsamples, fine roots were separated from the woody roots after air drying. We defined fine roots as those that disintegrated when rubbed between the fingers; woody roots did not. Total root surface area (for the fine roots) was calculated by assuming roots were cylindrical and multiplying total root length by π times diameter. Total number of mycorrhizal root tips was calculated by multiplying total number of root tips by percent mycorrhizal root tips.

Total-N and percent ¹⁵N enrichment for ground plant material were measured using the same procedure as for the glass fiber filters

used in the diffusions. Excess percent ^{15}N enrichment for the plant material was calculated as the difference between plant samples 48 h after the $^{15}\text{N-NH}_4^+$ application and plant samples (leaves) collected prior to injection. For all of the tree species, leaves, stems, woody roots and fine roots were separated and determination of carbon and nitrogen content made using a Europa ANCA-sl Elemental Analyzer (Europa Instruments, England).

Nitrogen immobilization, uptake, and mineralization calculations

To estimate the gross rate of mineralization and NH_4^+ consumption from the soil pool, we used the pool dilution calculations which accounted for the base level of ^{15}N in the NH_4^+ produced during mineralization (Blackburn 1979; Wessel and Tietema 1992). The base level of ^{15}N in the NH_4^+ was measured on the soil samples collected prior to the injection of ^{15}N . Average base levels of ^{15}N enrichment (%) in the soil NH_4^+ pool prior to injection was 0.391 ± 0.007 (mean \pm SE) for ambient and 0.399 ± 0.013 for elevated CO_2 (not statistically different). Pool dilution calculations were performed separately for the first and second 24 h following the injection of the ^{15}N label, as well as the full 48-h period. Each of these three sets (0–24 h, 24–48 h, and 0–48 h) of pool dilution calculations were performed using actual data on NH_4^+ pool sizes and ^{15}N content of the soil solution at the beginning and end of the time interval.

We assumed that all NH_4^+ consumption was due to plant and soil microbial uptake and therefore used the relative amount of ^{15}N tracer acquired by plants and soil microbes to derive NH_4^+ uptake rates by partitioning the pool dilution estimate of gross immobilization. As an additional check, we also estimated plant and microbial uptake using the total amount of ^{15}N tracer recovered in plant and microbial biomass. However, because the ^{15}N is a small and changing proportion of the pool of available NH_4^+ , recovery of ^{15}N in biomass will underestimate actual uptake rates (e.g., Barclaugh 1991; Davidson et al. 1991). We did not include nitrification rates when partitioning gross NH_4^+ consumption rates because standing pools of nitrate were extremely low (< 1 ppm in soil extracts) making measurement of the ^{15}N tracer in this pool not possible. While there is some data to suggest that gross nitrification rates can be high even when standing pools of nitrate in the soil solution are low (Davidson et al. 1992; Stark and Hart 1997), recent experiments have found that elevated gross nitrification rates are linked to elevated N inputs to the soil (Tietema 1997). Because we maintained our mesocosms with no external nitrogen inputs for 2 years and nitrate levels were so low, we assumed that nitrification rates were negligible. We focused measurements on NH_4^+ rather than NO_3^- because we were interested in quantifying the fate of newly produced inorganic nitrogen from plant-derived soil detritus, and NH_4^+ is the dominant form of inorganic nitrogen in the soils we were examining (Aber et al. 1993; Magill et al. 1997).

Statistical analysis

For all one-mesocosm, one-measurement variables, data were analyzed using a two-way ANOVA, with the CO_2 treatment and block as independent variables. For measurements made separately for each species or separately for plants and soil microbes, a three-way ANOVA was used with the addition of species/organism as an additional independent factor. These ANOVAs treated block as a fixed factor because individual CO_2 glasshouses were not a priori paired by CO_2 levels. For a given species, the effect of CO_2 was tested using linear contrasts (SuperANOVA, v1.11, Abacus Concepts, Berkeley, Calif.). Assumptions of homoscedasticity and normality of residuals were tested with normal probability plots (DataDesk v4.0, Data Description, Inc., Ithaca, N.Y.) and data were transformed (log or arcsine square root) in cases where it improved compliance with the assumptions of parametric statistics.

Results

Elevated CO_2 led to significant increases in net plant biomass production in the mesocosms (Table 1). Pooling above- and belowground biomass, CO_2 enhancement (average elevated CO_2 /average ambient CO_2) was 1.31. This is a level of net biomass enhancement close to the average level observed for most single plant studies (Ceulemans and Mousseau 1994; Poorter 1993; Wullschleger et al. 1995), and significantly greater than that observed in monocultures of yellow birch (Berntson and Bazzaz 1997; Wayne and Bazzaz 1995). The effect of elevated CO_2 on belowground biomass (62% increase) was greater than that for aboveground biomass (23%; Table 1). Total plant nitrogen content (above and belowground biomass for all species pooled together) was unaffected by elevated CO_2 , and thus C/N ratios of plant biomass were significantly increased (Table 1).

In total, we cataloged 13 different species regenerating within the mesocosms (trees: *Betula papyrifera*, *B. alleghaniensis*, *B. populifolia*, *Quercus rubra*; other: *Gaultheria procumbens*, *Rubus alleghaniensis*, *Carex* sp., *Medeola virginiana*, *Trientalis americana*, *Dennstaedtia punctilobula*, *Hieracium* sp., *Lysimachea quadrifolia*, *Panicum* sp.). Aside from a single mesocosm which was completely dominated by *Dennstaedtia*, by the second year every mesocosm was dominated by *B. papyrifera* and *B. alleghaniensis* ($> 95\%$ of total plant biomass). Except for the two dominant birch species and *G. procumbens*, no other species occurred in more than two

Table 1 Plant biomass, allocation patterns, molar C/N ratios for plants (all species pooled together) and soil microbes, and all measured pools of nitrogen. P = probability that null hypothesis is true (ambient CO_2 = elevated CO_2). (ER ratio of elevated CO_2 mean over ambient CO_2 mean)

	Units	Ambient CO_2		Elevated CO_2		P	ER
		Average	SE	Average	SE		
Plant biomass	g m^{-2}	893	42	1173	65	0.00	1.31
Root/shoot biomass	–	0.29	0.02	0.39	0.05	0.14	1.32
Aboveground biomass	g m^{-2}	695	38	853	49	0.08	1.23
Belowground biomass	g m^{-2}	198	12	320	34	0.01	1.62
Plant C/N	–	69.0	4.0	87.0	3.4	0.05	1.26
Soil microbial C/N	–	11.0	0.4	12.2	0.7	0.19	1.11
Plant nitrogen	g m^{-2}	10.6	0.6	10.9	0.8	0.77	1.03
Microbial N content	g m^{-2}	3.68	0.89	2.91	0.32	0.36	0.79
Extractable NH_4^+ -N	g m^{-2}	0.218	0.025	0.222	0.038	0.94	1.02

mesocosms per CO₂ level. This high level of rarity made it impossible to infer anything about changes in species diversity or richness in response to elevated CO₂. However, there was a significant relative increase in *B. alleghaniensis* and a significant relative decrease in *B. papyrifera* biomass in elevated CO₂ (Fig. 1). This significant change

in relative biomass contribution was driven by a significant absolute increase in *B. alleghaniensis* (Table 2A) biomass, and a non-significant change in absolute biomass in *B. papyrifera* (Table 2B) or any other species. The significant increase in *B. alleghaniensis* biomass was due to increases in the density as well as average size of

Table 2 A Summary of growth, allocation, nitrogen content, root architecture, mycorrhizal colonization and NH₄⁺ uptake rates for *B. alleghaniensis* and for *B. papyrifera*. *P* = probability that null

hypothesis is true (ambient CO₂ = elevated CO₂) (*ER* ratio of elevated CO₂ mean over ambient CO₂ mean)

<i>Betula alleghaniensis</i>	Units	Ambient CO ₂		Elevated CO ₂		<i>P</i>	<i>ER</i>
		Average	SE	Average	SE		
A							
Growth and allocation							
Plant density	m ⁻²	39.0	7.4	80.6	16.7	0.00	2.07
Average plant biomass	g plant ⁻¹	4.2	1.2	8.2	1.7	0.10	1.96
Total biomass	g m ⁻²	224	73	593	113	0.01	2.65
Root/shoot biomass	–	0.37	0.05	0.47	0.03	0.06	1.26
Fine root mass	g m ⁻²	13.0	4.7	46.3	8.9	0.00	3.57
Fine/woody root mass	–	0.37	0.04	0.37	0.04	0.95	0.99
Nitrogen content							
Whole plant nitrogen	g m ⁻²	2.98	0.99	4.82	0.64	0.18	1.62
C/N-leaves	–	21.4	1.0	28.8	1.7	0.00	1.35
C/N-stem	–	65.7	6.3	122.7	10.0	0.00	1.87
C/N-woody roots	–	67.9	6.4	111.8	9.8	0.00	1.65
C/N-fine roots	–	35.2	1.3	47.3	2.0	0.00	1.34
Root architecture & mycorrhizae							
Mycorrhizal colonization	% Tips	32.7	5.3	55.8	4.2	0.00	1.71
Root diameter	µm	162	3	154	4	0.14	0.95
Root length	km m ⁻²	2.95	1.08	12.44	2.24	0.00	4.22
Root surface area	m ⁻² m ⁻²	2.93	1.05	11.87	2.12	0.00	4.05
Specific root length	km g ⁻¹	0.23	0.02	0.27	0.02	0.09	1.19
NH ₄ ⁺ uptake rate							
¹⁵ N recovery ^a	mg ¹⁵ N m ⁻² day ⁻¹	6.9	3.6	4.3	0.8	0.22	0.61
Pool dilution 0–24 h ^b	mg N m ⁻² day ⁻¹	41.6	20.9	16.6	6.8	0.13	0.40
Pool dilution 24–48 h ^b	mg N m ⁻² day ⁻¹	17.6	12.2	1.9	4.7	0.30	0.11
Pool dilution 0–48 h ^b	mg N m ⁻² day ⁻¹	43.2	25.2	9.2	2.5	0.75	0.21
B							
<i>Betula papyrifera</i>							
Growth and allocation							
Plant density	m ⁻¹	64.6	8.0	43.6	6.9	0.02	0.68
Average plant biomass	g plant ⁻¹	11.3	1.6	11.0	2.3	0.86	0.97
Total shoot biomass	g m ⁻¹	658	70	593	116	0.69	0.90
Root/shoot biomass	–	0.25	0.01	0.35	0.05	0.05	1.42
Fine root mass	g m ⁻²	31.8	3.6	33.3	6.7	0.70	1.05
Fine/woody root mass	–	0.33	0.02	0.33	0.03	0.92	1.02
Nitrogen content							
Whole plant nitrogen	g m ⁻²	7.44	0.75	5.36	1.02	0.10	0.72
C/N-leaves	–	25.1	1.1	31.0	1.7	0.00	1.24
C/N-stem	–	115.6	3.7	118.3	1.2	0.73	1.02
C/N-woody roots	–	87.7	4.7	97.3	6.1	0.15	1.11
C/N-fine roots	–	38.3	1.8	48.0	2.0	0.00	1.25
Root architecture & mycorrhizae							
Mycorrhizal colonization	% Tips	36.3	5.2	52.7	2.3	0.00	1.45
Root diameter	µm	158	5	158	5	0.88	1.00
Root length	km m ⁻²	8.81	1.44	7.81	1.46	0.87	0.89
Root surface area	m ⁻² m ⁻²	8.5	1.3	7.57	1.4	0.87	0.89
Specific root length	km g ⁻¹	0.27	0.02	0.25	0.03	0.50	0.92
NH ₄ ⁺ uptake rate							
¹⁵ N recovery ^a	mg ¹⁵ N m ⁻² day ⁻¹	12.4	2.6	5.2	1.4	0.03	0.42
Pool dilution 0–24 h ^b	mg N m ⁻² day ⁻¹	81.7	26.5	11.1	29.2	0.07	0.14
Pool dilution 24–48 h ^b	mg N m ⁻² day ⁻¹	52.2	17.5	1.1	5.4	0.00	0.02
Pool dilution 0–48 h ^b	mg N m ⁻² day ⁻¹	65.7	17.9	14.1	6.3	0.00	0.21

^a Uncorrected uptake-total ¹⁵N recovered in plant biomass

^b Derived from partitioned pool dilution estimates of gross NH₄⁺ consumption

plants (Table 2A). For *B. papyrifera*, there was no significant increase in total biomass or average plant biomass, but there was a significant decrease in plant density (Table 2B).

Because the two dominant birch species comprised more than 95% of the biomass in all the mesocosms in

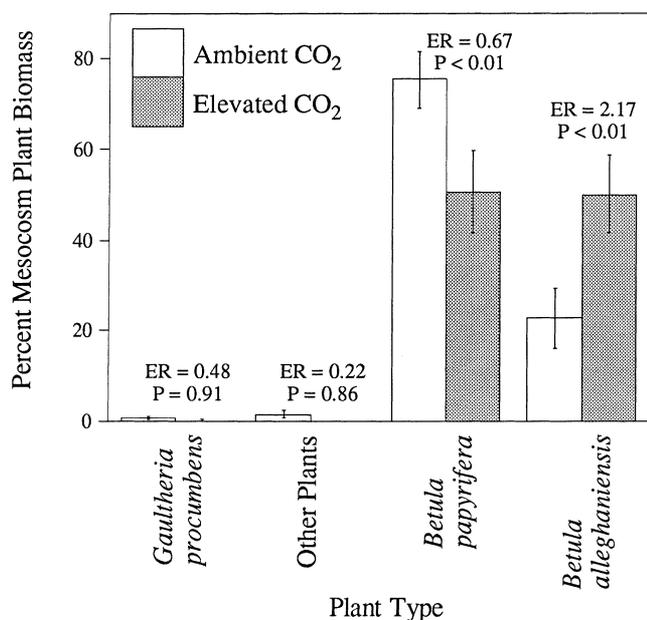


Fig. 1 Summary of the present of total biomass for each of the four groups of plants which were present in more than one mesocosm per CO₂ level. Percent biomass was calculated as the total biomass of a given plant group divided by the total plant biomass within the microcosm. The single soil monolith which was completed dominated by the fern was not included in this analysis (P probability that elevated CO₂ mean = ambient CO₂ mean, ER ratio of elevated CO₂ mean over ambient CO₂ mean). Error bars are 1 SEM

which they were present, we focused measurements of biomass allocation, root architecture, and mycorrhizal colonization rates on these species. In response to elevated CO₂ *B. alleghaniensis* showed significant increases ($P \leq 0.05$) in total biomass, fine root biomass, ectomycorrhizal colonization, fine root length, fine root surface area and specific root length; marginally significant increases ($P \leq 0.1$) in root/shoot ratios, and specific root length (SRL); and no significant change ($P > 0.1$) in total nitrogen content, the ratio of fine to woody root mass, or fine root diameter (Table 2A). In contrast, *B. papyrifera* showed significant increases only in root/shoot ratios and mycorrhizal colonization rates; and a marginally significant decrease in nitrogen content (Table 2B). Both species of *Betula* showed significant increases in C/N ratios for the leaves and fine roots in elevated CO₂ (Table 2A, B). For *B. papyrifera*, no significant changes in the C/N ratios of woody stems or roots were seen. In contrast, for *B. alleghaniensis* the C/N ratios of woody stems and roots increased significantly with elevated CO₂. We observed no significant change in NH₄⁺ uptake rates for *B. alleghaniensis*, regardless of the method used to estimate uptake rates (Table 2A). In contrast, we observed significant reductions in uptake rates of NH₄⁺ for *B. papyrifera* for every method we used to estimate uptake rates (Table 2B).

Observed rates of gross mineralization and NH₄⁺ consumption were strongly dependent on the time interval over which the flux rate was determined. On average, mineralization rates decreased by 71.5% and immobilization rates by 63.6% from the first to the second 24 h of the 48-h labeling period (Table 3). Estimates over the full 48 h incubation period were intermediate. Elevated CO₂ led to significant reductions in gross mineralization and consumption rates of NH₄⁺ for

Table 3 Results of pool dilution calculations, including gross mineralization and consumption, along with partitioned estimates of gross consumption (see Materials and methods for details) and ¹⁵N

recovery in plant and soil microbial biomass. P = probability that null hypothesis is true (ambient CO₂ = elevated CO₂) (ER ratio of elevated CO₂ mean over ambient CO₂ mean)

	Units	Time interval	Ambient CO ₂		Elevated CO ₂		P	ER
			Average	SE	Average	SE		
Gross mineralization rate	mg N m ⁻² day ⁻¹	0–24 h	242.0	64.0	189.0	90.0	0.56	0.78
		24–48 h	106.0	26.0	17.0	32.0	0.04	0.16
		0–48 h	201.2	41.1	103.7	25.0	0.06	0.52
Gross consumption rate	mg N m ⁻² day ⁻¹	0–24 h	262.0	66.0	173.0	89.0	0.40	0.66
		24–48 h	139.0	36.0	20.0	32.0	0.01	0.14
		0–48 h	215.3	45.7	101.1	28.1	0.05	0.47
Plant NH ₄ ⁺ uptake rate	mg N m ⁻² day ⁻¹	0–24 h	122.8	33.9	53.3	39.3	0.19	0.43
		24–48 h	69.6	19.8	7.2	9.1	0.00	0.10
		0–48 h	107.5	28.0	39.3	18.3	0.04	0.37
¹⁵ N recovery-plant ^a	%	–	35.5	7.1	20.6	4.8	0.10	0.58
Microbial NH ₄ ⁺ uptake rate	mg N m ⁻² day ⁻¹	0–24 h	138.9	49.8	119.9	58.4	0.78	0.86
		24–48 h	68.7	23.7	12.7	23.7	0.10	0.18
		0–48 h	107.8	35.1	61.7	15.6	0.23	0.57
¹⁵ N recovery-microbial ^a	%	–	30.5	7.8	39.7	5.0	0.35	1.30
Fraction plant ¹⁵ N ^b	%	–	55.7	8.2	32.9	4.7	0.02	0.59

^a Percent of total amount of ¹⁵N injected recovered in given pool

^b ¹⁵N acquisition by plants relative to the total amount recovered in plant and soil microbial biomass

the full 48 h incubation period, and for second 24 h, but not for the first 24 h.

Partitioned pool dilution estimates of plant NH_4^+ uptake rates were significantly reduced for the full 48 h incubation period and the second 24 h. However, we observed non-significant reductions in soil microbial uptake rates of NH_4^+ , though for the second 24 h of the incubation reduction in uptake was nearly significant ($P = 0.10$). The amount of ^{15}N recovered in plant and soil microbial biomass generally supported the pool dilution estimates. The decrease observed for plants was nearly significant ($P = 0.10$) but a non-significant change was observed for soil microbes. Expressed as a percentage of the total amount of ^{15}N recovered in plant and soil microbes, we observed a significant reduction in the amount of ^{15}N recovered by plants relative to soil microbes.

Discussion

Species composition

Species composition plays an important role in determining terrestrial ecosystem structure and function (Bassow 1995; Clark 1993; Pastor and Post 1986). Changes in composition of regenerating forests in an elevated CO_2 environment may itself represent a mechanism of long-term shifts in ecosystem productivity and nutrient cycling. The effects of elevated CO_2 on early tree growth and establishment may be a critical determinant of future forest composition. For example, assuming that the effect of elevated CO_2 on growth enhancement for several tree species is transitory, Bolker et al. (1995) used a forest simulator (Pacala et al. 1993) to show that elevated CO_2 may potentially lead to large changes in the composition and productivity of temperate forests.

In this study, we observed a significant shift in relative dominance of the two most abundant tree species regenerating within the mesocosms. In elevated- CO_2 environments, yellow birch (*B. alleghaniensis*) increased in frequency and white birch (*B. papyrifera*) decreased in frequency. Trees in the genus *Betula* produce large numbers of small seeds, typically have fast seedling growth rates, and often dominate early successional forest communities (Burns and Honkala 1990). These characteristics explain why the two dominant regenerating species were *Betula*. A review of the average growth enhancement of *B. alleghaniensis* and *B. papyrifera* in elevated CO_2 from individually grown plants does not *a priori* suggest that either species would benefit more than the other in response to elevated CO_2 . Net growth enhancement in a doubled- CO_2 atmosphere for individually grown *B. alleghaniensis* range between 23 and 86% (Rochefort and Bazzaz 1992; Wayne and Bazzaz 1995) and for *B. papyrifera* between 24 and 100% (Bazzaz et al. 1990; Berntson and Bazzaz 1996b; Rochefort and Bazzaz 1992). Thus, the increased rep-

resentation of *B. alleghaniensis* which we observed could not have been predicted on the basis of average individual-plant CO_2 growth enhancements. This discrepancy between actual changes in community composition and predicted changes is a critical challenge in predicting community and ecosystem effects of elevated CO_2 (Bazzaz and McConnaughay 1992; Körner 1996).

Shifts in the dominance from *B. papyrifera* to *B. alleghaniensis* in regenerating forest ecosystems in the Northeastern U.S. in response to elevated CO_2 is likely to be the result of an increase in water use efficiency in elevated CO_2 (Field et al. 1995). *B. alleghaniensis* typically regenerates in much moister microsites than does *B. papyrifera* (Burns and Honkala 1990). We observed a 13% increase in gravimetric moisture content with elevated CO_2 , and this may be an important factor leading to the differential performance of these species. It is not clear what the long term implications of such a shift would be. *B. alleghaniensis* is a longer lived, much larger tree than is *B. papyrifera*. However, communities dominated by *B. alleghaniensis* may be more sensitive to periods of drought.

Root architecture and mycorrhizal colonization

Along with effects on biomass production and allocation, elevated CO_2 levels can lead to changes in root architecture, physiology, and mycorrhizal symbiosis. In general, elevated CO_2 appears to lead to changes in root form and function which lead to increases in potential nutrient acquisition (Berntson and Bazzaz 1996a; Norby 1994; Rogers et al. 1994). Changes in plant root system architecture in elevated- CO_2 environments include greater total root length, an increase in the amount of soil explored, and more root tips (Berntson and Woodward 1992; Kaushal et al. 1989; Larigauderie et al. 1994; Rogers et al. 1992). In the majority of studies to date, elevated CO_2 has led to significant increases in the colonization of trees by ectomycorrhizal fungi (Berntson et al. 1997; Godbold and Berntson 1997; Ineichen et al. 1995; O'Neill 1994). Together, these changes in root architecture and mycorrhizal symbiosis have the potential of increasing the total amount of nutrients a plant can acquire from the soil (Berntson 1994; Yanai et al. 1995).

In this study we found that elevated CO_2 led to a significant increase in the overall size of the root systems of *B. alleghaniensis* but not *B. papyrifera*, as measured by fine-root length, number of root tips, and root surface area. This differential belowground response between the two dominant tree species was not due to enhanced allocation of biomass belowground, as *B. papyrifera* showed a greater enhancement in root/shoot biomass than did *B. alleghaniensis* in elevated CO_2 (Table 2A, B). Differences in root system size between these species is largely related to plant size (e.g., biomass). In contrast to root architecture, elevated CO_2 led to increases in the degree of ectomycorrhizal colo-

nization for every tree species in the experimental mesocosms. Taken together, the changes in root architecture and mycorrhizal symbioses imply that elevated CO_2 leads to an increase in the potential of regenerating trees to acquire nutrients from the soil.

Plant productivity and plant C/N

In this study, elevated atmospheric CO_2 resulted in a 31% increase in plant biomass (pooling all plant species, above and belowground biomass), without a significant change in total plant nitrogen content (Table 1). This imbalance in biomass production relative to nitrogen acquisition led to a 26% increase in whole-plant C/N ratios. From the perspective of plant growth, increased C/N ratios represent an increase in nutrient use efficiency (Sheriff et al. 1995; Vitousek 1982), allowing plants to maintain or increase biomass production with limited N.

However, from the perspective of decomposing plant detritus, increased C/N ratios of plant biomass in elevated CO_2 has important implications for the long term storage of biomass in terrestrial ecosystems as well as potential feedbacks on nitrogen cycling and nitrogen availability (Rastetter et al. 1992; Shaver et al. 1992). Studies of decomposition rates in the field have shown that higher C/N ratios can lead to slower decomposition rates and increased nitrogen immobilization (Aber et al. 1990; Downs et al. 1996; Melillo et al. 1982). However, it has not been conclusively demonstrated that atmospheric CO_2 -induced changes in plant C/N ratios will lead to alterations in the rate of plant detritus decomposition. Some studies have observed reduced decomposition (mass-loss) rates of plant detritus from elevated CO_2 environments which have elevated C/N ratios (Boerner and Rebeck 1995; Cotrufo and Ineson 1995). Other studies have found that reductions in decomposition rates can be either insignificant or transitory (Coûteaux et al. 1991; O'Neill and Norby 1996).

The study we present here was carried out for just under 2 years. At the end of this time, we found no effect of elevated CO_2 on total plant-N content. Thus, we would not expect any N limitations due to the sequestration of N in plant biomass or detritus for early stages of forest regeneration. Instead, we suggest the relationship between plant C/N ratios and mineralization of plant-derived detritus may be a more important feedback on nitrogen availability, and thus possibly plant growth responses to elevated CO_2 . To assess the potential for changes in plant C/N ratios to influence decomposition rates in the context of this study, it is important that we examine those plant organs which turnover rapidly and therefore represent the sources of plant-derived detritus in the soil (e.g., leaves and fine roots in this study). The C/N ratios of leaves and fine roots were significantly increased in elevated CO_2 (by 24–35%) for both of the dominant tree species.

The interplay between nitrogen cycling and plant capacity for nitrogen uptake

We used pool dilution techniques to quantify *in situ* gross mineralization and immobilization rates of nitrogen within the soil. Pool dilution techniques are more useful for characterizing potential changes in nutrient availability than the dynamics of mass loss because they directly quantify the fluxes, and thus potential availability, of nitrogen. The dynamics of mass loss do not provide a valid estimate of nitrogen mineralization because plant detritus can immobilize a significant amount of nitrogen during early decomposition phases, especially when C/N ratios are high (Aber and Melillo 1991; Aber et al. 1990; Downs et al. 1996). This disparity between mass and nitrogen dynamics in decaying litter suggests that to determine potential effects of altered plant detritus quantity and/or quality on nitrogen availability within the soil we need to directly quantify fluxes of nitrogen.

Using pool dilution calculations, we observed reductions in NH_4^+ mineralization and consumption rates under elevated CO_2 (Table 3). These results are consistent with and possibly the result of the increases in C/N ratios for the fine roots and leaves. These results do not support either of the feedback hypotheses presented by Zak et al. (1993) or Díaz et al. (1993). We found no evidence to support the suggestion of Zak et al. (1993) that elevated CO_2 would lead to increased nitrogen availability. Rather, we observed significant reductions in gross mineralization and consumption rates of NH_4^+ in elevated CO_2 . The results of this study only partially support the suggestion of Díaz et al. (1993) that increased nitrogen immobilization by soil microbes for inorganic nitrogen will reduce plant available N. We found that NH_4^+ uptake was reduced for plants, but not clearly so for soil microbes. In terms of the relative amount of ^{15}N tracer recovered in plant and soil microbial biomass, there was a shift from what appeared to be a balance in ambient CO_2 (a 50/50 split in acquired tracer) to a significant reduction in the relative amount of tracer acquired by plants in elevated CO_2 (Table 3). However, neither total microbial-N content nor total plant-N content were significantly affected by elevated CO_2 (Table 1). The results we present here are more consistent with a previous study we carried out examining the growth and nitrogen dynamics in monocultures of *B. alleghaniensis* (Berntson and Bazzaz 1997) than with either Zak et al. (1993) or Diaz et al. (1993).

A critically important clarification to make regarding the data we present is that all the data on estimated fluxes (mineralization, consumption, and uptake of NH_4^+) correspond to the last 48 h of the experiment, while all of the data on carbon and nitrogen pool sizes integrate processes over a much longer period of time. This clarification is important because our conclusion regarding the presence or absence of a negative feedback on potential plant productivity through nitrogen avail-

ability will vary depending on which set of data we examine.

Judging by total nitrogen content measured in plant and soil microbial biomass at the end of the experiment, elevated CO₂ had no effect on net nitrogen acquisition for either group of organisms, nor on their relative ability to acquire nitrogen (Table 1). The effect of elevated CO₂ on nitrogen mineralization, consumption, and uptake rates which we quantified at the end of the experiment lead to a very different conclusion regarding the potential impact of elevated CO₂ on nitrogen cycling and nitrogen availability to plants. The large, significant reductions in nitrogen cycling (production and consumption of NH₄⁺) and plant uptake we observed, if extrapolated into the future, suggest that reduced plant availability of nitrogen is likely to constrain positive growth enhancements in elevated CO₂ due to nitrogen limitation.

The key question which follows from these apparently contrasting data is "How is it possible for these data to be so different from each other?" One possible answer to this question is simply that rates of nitrogen production and consumption were not constant throughout the duration of the experiment. This hypothesis, while not directly testable with our data, could explain the apparent discrepancy in pool sizes of nitrogen and measured flux rates. If the early effects of elevated CO₂ on plant growth were increases in plant size and plant ability to acquire available nitrogen (though changes in root architecture, mycorrhizal symbioses, and possibly root physiology), there may have been an early enhancement in plant growth and nitrogen uptake. This a reasonable assumption, as elevated CO₂ typically leads to larger growth enhancements early in plant ontogeny (Bazzaz et al. 1993). Further, when the mesocosms were first established, all of the nitrogen supplied within the soil via mineralization was derived from labile organic matter within the soil. Presumably, the different mesocosms had about the same amount of nitrogen available because they were all watered the same amount and maintained in the same temperature regimes. If, through time, the production of high C/N detritus in elevated CO₂ led to a reduction in NH₄⁺ mineralization rates and thus consumption rates, the early increases in growth and nitrogen acquisition would diminish. This scenario is consistent with all of the data presented in this study. However, to critically test this hypothesis, future studies must examine the ontogeny of nitrogen cycling and nitrogen uptake in plant communities.

One additional important factor which this study has demonstrated is that even though we observed changes in root architecture (root length, root surface area; particularly in *B. alleghaniensis*) and ectomycorrhizal colonization which would, all other factors being equal, increase the ability of plants to acquire nitrogen, we observed no increase in plant uptake of NH₄⁺. There are two possible hypotheses for this pattern. First, that changes in root architecture and mycorrhizal colonization are not important in determining plant ability to

acquire soil nutrients (NH₄⁺ in this case). Second, that the reduction in the rate of supply of NH₄⁺ led to a constraint in nitrogen availability which was more important than the observed changes in root growth. We suggest that the first hypothesis is not consistent with our data. In *B. alleghaniensis* we observed a significant increases in root architectural parameters and mycorrhizal colonization (Table 2A), and no statistically significant drop in total ¹⁵N acquisition or estimated NH₄⁺ uptake. In contrast, in *B. papyrifera* we observed no change in root architectural parameters (though we did see an increase in mycorrhizal colonization; Table 2B) and we observed a decrease in both total ¹⁵N acquisition or estimated NH₄⁺ uptake. These data suggest that changes in root architecture do play an important role in mediating plant ability to acquire nitrogen, even when total plant uptake is reduced to reductions in availability due to environmental limitations.

Acknowledgements Special thanks to P. Wayne for help collecting the soil monoliths, J. Nevins and J. McCarthy for access to the Europa, A. Li, R. Keane and C. Stodolski for invaluable help in harvesting and processing the plant samples, and M. Jasienski, S. Catovsky, W. Bossert, M. T. Sebastia Alvarez and four anonymous reviewers for helpful comments on the manuscript. This research was supported by a U.S. D.O.E. Graduate Fellowship in Global Change and a U.S. N.O.A.A. Postdoctoral Fellowship in Global Change to G.M.B and a grant from N.I.G.E.C. to F.A.B.

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