

Carbon cycling across ecosystem succession in a north temperate forest: controls and management implications

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doi.org/10.6073/pasta/b47e7e6f05d6fcd7136c1e28f23e5217), and flux tower data (Gough et al. 2023a; 2023b; doi.org/10.17190/AMF/2204882; doi.org/10.17190/AMF/1881597). Additional datasets will be published in an EDI repository upon manuscript acceptance.

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Abstract

Despite decades of progress, much remains unknown about successional trajectories of carbon (C) cycling in north temperate forests. Drivers and mechanisms of these changes, including the role of different types of disturbances, are particularly elusive. To address this gap, we synthesized decades of data from experimental chronosequences and long-term monitoring at a well-studied, regionally representative field site in northern Michigan, USA. Our study provides a comprehensive assessment of changes in above- and belowground ecosystem components over two centuries of succession, links temporal trends in major C pools and fluxes with underlying drivers, and offers several conceptual insights to the field of forest ecology. The first of these concepts is that temporal dynamics in several ecosystem components are consistent across severe disturbances that reset succession and partial disturbances that slightly modify it. Namely, both types of disturbance increase soil N availability, alter fungal community composition, and affect aboveground growth and interspecific competition as short-lived pioneer trees succeed to longer-

lived taxa. These changes in turn affect soil C stocks, respiratory emissions, and other belowground processes. In contrast, other ecosystem components have inconsistent effects on C cycling over succession. For example, canopy structure does not influence C uptake early in succession, but becomes important as stands develop, with the importance of individual structural properties changing over the course of two centuries of stand development. In recent decades, we find that climate change is masking or overriding effects of community compositional change on C uptake; respiratory emissions appear sensitive to both climatic and successional compositional change. In synthesis, we emphasize that time is not a driver of C cycling, it is a dimension within which ecosystem drivers such as canopy structure, tree and microbial community composition change. Changes in those drivers, not in forest age, are what control forest C trajectories, and those changes can happen quickly or slowly, through natural processes or deliberate intervention. Stemming from this view and a whole-ecosystem perspective on forest succession, we offer management applications from this work and assess its broader relevance to understanding long-term change in other north temperate forest ecosystems.

Key words: forest disturbance, ecosystem development, biogeochemistry, community composition, climate change

1. Introduction

The general trajectories of carbon (C) stocks and sequestration rates over the course of secondary succession in north temperate forests are well known. Rapid sequestration during the initial decades after disturbance begins to decline later in the first century of stand development, while standing C stocks continue to increase, potentially for centuries (Pan et al. 2011; Pregitzer and Euskirchen 2004). However, there is considerable variation in these trends, evidenced by large amounts of variance in age-C relationships within individual sites, sites that show the same general patterns over very different timescales, and sites that depart from general trends altogether (Birdsey et al. 2023; Bradford and Kastendick 2011; Keeton et al. 2011; Thom and Keeton 2019). This variation indicates that our understanding of the factors controlling forest C trajectories has room to grow; it also highlights the fact that forest age is far from the only thing that influences C cycling over decade to century timescales.

Because all approaches to studying ecosystem succession struggle with its long timescales, sites that combine multiple complementary approaches can make powerful contributions to our understanding of the topic. Chronosequences have limitations, but as yet they are the only way to empirically quantify C pools and fluxes over centuries (Wardle et al. 2012; Yanai et al. 2003).

The other empirical approach—long-term monitoring—allows for longitudinal observation and evaluation of trends inferred through chronosequences, but even the longest-running of such studies cannot capture the centuries-long process of ecosystem succession (Eisen and Barker Plotkin 2015; Fahey et al. 2005; Hoover et al. 2011; Novick et al. 2018). In combination, these two complementary approaches can identify patterns and mechanisms of successional change, but even still it can be challenging to place results in the context of north temperate forests more

broadly. Such context, however, is necessary to produce transferable insights based on mechanistic understanding.

Long-term studies indicate several factors likely to control C stocks and sequestration rates over successional timescales in north temperate forests. The importance of forest structure—particularly canopy structure—to C cycling has often been demonstrated, though typically not through a successional lens (Gough et al. 2019; Ishii et al 2004; Reich 2012). Where it has been investigated as an ecosystem property that develops over the course of succession, increasing canopy structural complexity has been shown to increase forest C uptake by increasing resource use efficiency (Gough et al. 2022; Hardiman et al. 2013; Pan et al. 2013; Scheuermann et al. 2018). However, canopy structural complexity is to some extent linked to another important driver of forest C: forest community composition (Pedro et al. 2017). Trees are the most readily observed aspect of the forest community, and their compositional change over time is used almost synonymously with succession (although succession is a whole-ecosystem process). As a driver, the composition of the tree community determines canopy functional traits that influence many ecosystem processes, including C cycling (Fahey et al. 2019). Tree community composition also affects forest C in other ways, including interspecific variation in hydraulic strategies, wood density, aboveground vs. belowground C allocation, and interactions with the composition of other organismal communities, such as soil microbial communities (Abramoff and Finzi 2016; Davidson et al. 2002; Prescott and Grayston 2013, Woodcock and Shier 2002). Soil microbial communities in turn connect to a third major driver of forest C pools and fluxes: soil biogeochemistry. Soil biogeochemistry in this sense encompasses successional changes in soil properties such as pH (Compton et al. 1998; Flinn and Marks 2007), as well as microbial communities and the C- and nutrient-transforming processes that they mediate (Goodale and

Aber 2001; Knelman et al. 2017; Lewis et al. 2014; Lovett et al. 2018). Nitrogen (N) is particularly important among nutrients because it limits primary production, and the forms of it that are available for plant and microbial uptake change over the course of succession (Leduc and Rothstein 2010; Nave et al. 2009). Moreover, different microbial taxa play different roles in mineralizing organic matter and releasing soil N (Phillips et al. 2013). In acidic northern forest soils, ectomycorrhizal (EcM) fungi are major players in N cycling, and EcM taxa that explore larger volumes of soil or produce organic matter depolymerizing enzymes play more prominent roles as succession advances and a larger share of soil N pool is held in less accessible organic forms (Chen et al. 2019; Hobbie et al. 2013; Leduc et al. 2013). Successional interactions between soil N availability, photobiont C supply, and mycobiont enzyme production and N foraging are as yet better characterized in boreal forest soils, but are relevant to the N economy and C balance of temperate forest soils as well (Baldrian et al. 2023; Fosmark et al. 2024; Hay et al. 2015; Jorgensen et al. 2022). Research that places these biogeochemical processes in the context of aboveground ecosystem dynamics (e.g., forest composition, canopy structure) is particularly needed to better understand succession as a whole-ecosystem process.

In working towards the goal of transferable, predictive science, enumerating the factors that influence forest C over succession is arguably less important than describing how these factors operate holistically in ecosystems. Inspired by recent multi-decadal C cycling assessments from intensively studied north temperate forests (Desai et al. 2022; Finzi et al 2020; Hollinger et al. 2021), we synthesize new data and past results from long-term monitoring and chronosequence experiments at a well-studied site, the University of Michigan Biological Station (UMBS), in northern Lower Michigan. Our analysis is intended to extend this body of work by investigating longer-term successional changes spanning decades to centuries, by: (1) identifying trends and

drivers of major C pools and fluxes; (2) describing mechanisms by which they operate; (3) assessing relevance to the ecology and management of north temperate forests. The first two of these objectives are facilitated by offering several predictions that inform the C pools, fluxes, and drivers we have elected to focus on here. First, we predict that successional development of forest structure regulates vegetation growth and overall ecosystem C accumulation. Second, we predict that changes in aboveground (tree) and belowground (microbial) community composition are coupled through successional time, with corresponding influences on biogeochemical functioning. Third, we predict that concurrent changes in composition and environmental drivers have influenced trends in ecosystem C uptake and loss over recent decades.

2. Methods

2.1 Study site- UMBS (45.56°, -84.72°) is in northern Lower Michigan, in the southeastern 1/3 of the 261,000 km² Laurentian Mixed Forest Province of the USDA Forest Service ecosystem classification framework (Cleland et al. 1997; McNab et al. 2007). The province has a humid continental climate with cold winters and warm summers; at UMBS, mean annual temperature (MAT; 6.9°) and mean annual precipitation (MAP; 893 mm, including 284 cm snowfall; 1991-2020 for all parameters) fosters mixtures of warmer-temperate (*e.g.*, oak, maple) and cold-tolerant boreal (*e.g.*, spruce, fir, birch) forest types. Land use and disturbance history is broadly consistent throughout the province, where millennia of Native American ecosystem management activities were displaced by clearcutting and uncontrolled wildfires beginning with Euro-American colonization in the late 1800s and extending into the early 1900s (Lorimer 2001; Rhemtulla et al. 2009; Whitney 1987). Modern forest management began around the middle of the 20th century, and is more extensive than intensive: early-successional deciduous and mixed forests are clearcut and regrown over 40-80 year rotations, longer-lived deciduous cover types

experience selection harvests at 10-20 year intervals, and upland conifers (mostly pines) are managed with periodic thinning and regeneration harvests at 60-100 years of age (Bates et al. 1993; Gahagan et al. 2015; Gerlach et al. 2002; Palik et al. 2003; Stone 2002).

A detailed description of the landscape setting and experimental design of UMBS (Nave et al. 2017a) is summarized here. The regional Silurian and Devonian sedimentary bedrock lays buried beneath 100-200 m of glacial and lacustrine sediments deposited during the late Pleistocene. The highest elevations on the landscape are moraines of poorly-sorted till deposited directly by the continental ice mass; these are exceeded in area by lower-lying outwash plains, which were deposited by meltwater flowing away from ice margins. These landforms were modified during regional glacial re-advances 12,600 -10,500 years before present, and, at the lowest elevations, by lacustrine processes 4,300 - 3,000 years before present (Schaetzl et al. 2002; Nave et al. 2017b). Soils formed in till are mostly coarse-loamy Lamellic and Alfic Haplorthods (USDA Subgroups), outwash soils are predominantly sandy Entic Haplorthods, and soils in the lowest positions are predominantly Endoaquods, Endoaquents, and Haplosaprists (USDA Great Groups) formed in lake-modified outwash sediments and Holocene organic deposits.

Current vegetation at UMBS is the local result of the land use history that occurred across the wider ecological province. Namely, clearcutting and wildfires between 1870-1923 replaced pre-exploitation forests of long-lived species including red pine (*Pinus resinosa*), white pine (*P. strobus*), eastern hemlock (*Tsuga canadensis*), sugar maple (*Acer saccharum*), American beech (*Fagus grandifolia*), and red oak (*Quercus rubra*) with mixed deciduous-conifer forests dominated by early-successional taxa such as bigtooth aspen (*Populus grandidentata*) and paper birch (*Betula papyrifera*) in xeric to mesic settings, and trembling aspen (*P. tremuloides*) and balsam fir (*Abies balsamea*) in mesic to hydric settings. Most UMBS forestland is dominated by

early-successional taxa, which have been increasingly replaced by longer-lived species in the last 20-30 years, due to lack of disturbances or management (Gough et al. 2010). Isolated portions of UMBS property were cut partially, or not at all, or were burned before cutting could occur and therefore serve as older reference forests, with composition and structure that might be more extensive today if not for the widespread disturbances of a century ago. Upland forests at UMBS are broadly representative of the wider Laurentian Mixed Forest Province in terms of their composition, growth rates, and disturbance history (Gough et al. 2007; Nave et al. 2017a).

2.2 Experimental design- This study is based upon data collected through 2020 from several long-term experimental designs at UMBS (Fig. 1), with unique aspects detailed below.

2.2.1. Chronosequences and old reference forests- Two experimental chronosequences, differing in the type of stand-replacing disturbance, allow for the observation of stands ranging in age from 22 – 109 years. All chronosequence stands are on sandy outwash plains and have similar soil properties. Individual chronosequence stands were established either by clearcut + residue burning (known as the Burn Plots; Scheiner and Teeri 1981), or by a clearcut not followed by an intentional residue fire. Given that nearly all of the UMBS landscape (and the region) experienced clearcutting and fire ~100 years ago, the two chronosequences therefore differ in whether they have experienced one stand-replacing cut-and-burn disturbance (hereafter 1x cut+burn) or two stand-replacing cut-and-burn disturbances (2x cut+burn). In 2020, stand ages in the 1x cut+burn chronosequence were 33, 48, 68, and 109 years old (reflecting clearcutting in 1987, 1972, 1952, and 1911, respectively); 2x cut+burn stands were 22, 40, 66, 72, and 84 years old (clearcut + residue burning in 1998, 1980, 1954, 1948, and 1936, respectively).

The experimental chronosequences are complemented by observations of older reference stands, some of which have been monitored since the 1940s (Fig. 1). The oldest of the three reference

stands reported here, Indian Point, is a hemlock-northern hardwoods forest with a long history of Native American management (Albert and Minc 1987); it is uneven-aged, but in the area that has been intensively monitored, dominant trees date to *ca.* 1830 (i.e., stand age = 190 years). The two other old reference stands are pine-dominated, with canopy dominants dating to 1885 – 1890 (stand age = 130 years). One of these stands was partially cut and lightly burned; the best available records (and site evidence) suggest the other was initiated after a stand-replacing fire. Due to their unique disturbance histories and differences in soils, landforms, and topoclimate, none of the three reference stands is an ideal representation of conditions that would be expected on the paired chronosequences after additional decades of development (Nave et al. 2019).

In both the experimental chronosequences and the old reference stands, individual stands range from just over 1 ha to more than 10 ha in size; in each stand, most data collection occurs within 3 plots, each 0.1 ha in size (except for 2 plots, 0.07 and 0.14 ha, in the 22-year-old “Burn Plot”). Throughout the remainder of this paper, we use the term “stand” to describe an area of forest of a known common age, composition, and disturbance, from one to several hectares in size. We distinguish stands from plots, which are sampling units of logistical convenience, but note that in many stands, individual plots were arrayed to sample across measurable or visually apparent spatial variation in ecosystem structure, disturbance severity, or to minimize edge effects. The sole exception to the “stand” vs. “plot” terminology is the Burn Plots, which by this definition consist of stands differing in ages, but whose name we reference (with capitalization) owing to their immediate recognition for generations of UMBS researchers.

2.2.2 Flux towers- The other experimental design that we draw upon for this analysis consists of paired reference and treatment flux towers known in the AmeriFlux Network as the US-UMB (reference) and US-UMd (disturbed) core sites (Fig. 1). The reference tower has a ~100 ha

footprint and is supported by ground-based monitoring of ecosystem properties, C pools and fluxes across an array of 81 permanent plots since 1997. The treatment tower, erected in 2006, has a ~30 ha footprint where all of the aging, early-successional trees (aspen and birch) were stem girdled in 2008 to accelerate an incipient successional transition to longer-lived taxa- a process that has continued to unfold in the reference footprint since that time. The treatment footprint holds 22 permanent sampling plots. In each footprint, a single 1.1 ha plot surrounds the flux tower, and the remainder of the plots (0.08 ha each) are located at 100 m intervals along transects that radiate away from the tower. Within the scientific literature, comparisons of ecosystem processes and C cycling between the reference and treatment footprints have occasionally been reported as results of the Forest Accelerated Succession Experiment (FASET; Nave et al. 2011). In this paper, we refer to the reference vs. treatment towers explicitly when differentiating between slower natural background succession vs. experimentally accelerated succession, or to FASET (as a paired design) when drawing inferences from the two footprints collectively. Like the paired chronosequences, the footprints of both flux towers are located principally on sandy, high-level outwash plains.

2.3 Data collection methods- This paper synthesizes existing and presents previously unpublished data collected over the last 40 years on the experimental designs described in section 2.2. The following subsections provide overview descriptions; details are provided in references cited here and in Appendix S1.

2.3.1 Ecosystem C stocks- Inventories supporting the estimation of aboveground live tree, standing dead tree, and coarse down woody debris C stocks are typically completed at 5-year intervals in the experimental designs reported in this paper. At each census, permanent plots are inventoried for all trees ≥ 1.37 m in height, which are identified to species and measured for

diameter at breast height. Aboveground biomass C stocks are calculated from these inventory data using a combination of site-specific and generalized allometric equations (Nave et al. 2017a) and appropriate plot area expansion factors. For presentation and discussion, live trees are grouped into short-lived deciduous (aspen, paper birch), long-lived deciduous (all other deciduous taxa), and long-lived coniferous functional groups (all coniferous species found in these plots are long-lived). Coarse down woody debris (DWD) is censused for dimensions, density, and species on 3 subplots per 0.1 ha permanent plot, and scaled up using expansion factors (Clay et al. 2022). Soil, coarse and fine root C stocks were inventoried in 2014 to a depth of 1 m, as described in Nave et al. (2019). All data are publicly available (Nave et al. 2024).

2.3.2 Canopy structure- In this paper we report canopy structural parameters collected using ground-based portable canopy LiDAR in the individual plots located in each chronosequence and old reference forest stand. These parameters were collected along two perpendicular transects (40 m each) in each plot (Fahey et al. 2019; Scheuermann et al. 2018). From the raw LiDAR returns, we used the *forestr* package to compute three canopy structural parameters for analysis, including leaf area index (LAI), mean maximum canopy height, and canopy rugosity (Hardiman et al. 2013; Atkins et al. 2018). These canopy structural data have been previously published (Atkins 2002; Fahey et al. 2019; Scheuermann et al. 2018), but not used to address the specific questions of the present synthesis.

2.3.3 Biometric C flux measurements- This paper reports four major C fluxes derived from plot-based biometric measurements: wood net primary production (NPP_w), soil respiration (R_s), fine root production, and leaf litterfall. Wood NPP was calculated for all experimental chronosequence and old reference forest stands as the change in woody biomass C between 2019 and 2014 census intervals, divided by the 5-year remeasurement interval. Soil respiration

measurements were made on *in situ* collars across a range of dates and soil temperatures each year in the experimental chronosequence and old reference forest stands from 2014-2021, and from 1998-2020 and 2008-2017 in the UMB and UMd tower footprints, respectively. Detailed description of data collection and scaling to stand-level means is available in Clay et al. (2022) and Appendix S1. Fine root production, measured in chronosequence and old reference stands during the 2016-2019 growing seasons, is also described in Clay et al. (2022). Briefly, we used plastic mesh cores packed with freshly sieved, root-free native soil in each stand to quantify growing season root production (5-6 month) each year, averaged across all years within each stand. Soil respiration and fine root production data are publicly available (Nave et al. 2024).

Leaf litterfall fluxes reported in this paper come from a temporally consistent subset of the 0.08 ha permanent plots in the reference (n=12, 1997-2019) and treatment (n=14, 2006-2019) flux tower footprints. In each plot, the contents of individual traps (n=3) were aggregated into plot-level total leaf litterfall, which was sorted by species and component, and used to calculate annual means according to Gough et al. (2008).

2.3.4 Tower-based CO₂ and water vapor fluxes- In this paper, we report annual values of gross ecosystem C uptake (as gross primary production; GPP), gross ecosystem C emission (as ecosystem respiration; R_{eco}), and net ecosystem C uptake (as net ecosystem exchange; NEE) from publicly available AmeriFlux data for the US-UMB (2000-2020) and US-UMd (2008-2020) towers using the AmeriFlux-FLUXNET processed version of the data (Gough et al 2023a; 2023b). Annual values have been screened, gap-filled, and computed from the raw flux data collected on the two towers according to FLUXNET methods described in Pastorello et al. (2020). We also present growing season, midday (1100 – 1400 local time) ecosystem water use

efficiency (WUE) for reference (2001-2020) and treatment (2007-2020) towers (calculated from the same AmeriFlux-FLUXNET data sources), computed only for non-gap-filled periods.

2.3.5 Soil C dynamics- We examine changes in soil C using two distinct datasets and approaches to address two distinct questions. First, to assess soil C trajectories over a century of stand development, we compiled data from five known sampling campaigns on the Burn Plots. Across these campaigns, only O horizons were sampled consistently. Assuming differences in individual judgment and specific sampling techniques, we used unweighted effect-size meta-analysis to quantify the magnitude and temporal patterns of O horizon C stocks following cut+burn disturbance. Statistical methods follow Nave et al. (2021). Second, to assess how soil C stocks have changed in recent decades, we report results from a longitudinal re-sampling of an unmanipulated reference area adjacent to the Burn Plots. This area of 1911-origin (clearcut + unintentional residue fire) forestland was sampled in 1980, and again in 2009, providing an opportunity to compare mineral soil C stocks by genetic horizon over a 29-year period, independent of any disturbances other than the original region-wide cut+burn disturbance.

2.3.6 Soil biogeochemistry and microbial communities- In this paper, we report results from a systematic 2015 biogeochemical and microbial characterization of Burn Plots topsoils (A horizons). These samples allowed for comparison of fungal communities, extracellular enzyme activity, root biomass, microbial biomass C, soil N availability, soil C and N concentrations, pH, and soil moisture across the 5 experimental stands comprising the 2x cut+burn Burn Plots (data published in Nave et al. 2024). All preparation and most analysis steps were completed in the UMBS Analytical Laboratory, with sequencing at the Microbial Systems Molecular Biology Laboratory at the University of Michigan. Detailed methods are provided in Appendix S1. To inform fungal N cycling aspects of these measurements, we collected fungal sporocarps in 2022

from paired plots (n=6 plots) in the reference and treatment flux tower footprints that received ^{15}N tracer additions in 2010 (Nave et al. 2013). We analyzed these fungal samples by isotope ratio mass spectrometry in the UMBS Analytical Laboratory in order to assess long-term N foraging of several ectomycorrhizal (EcM) fungal taxa, using ^{15}N -enrichment as an indicator.

2.3.7 Climatology and atmospheric deposition- Daily climatology and weekly atmospheric deposition have been monitored since the late 1970s on the main campus of UMBS on the south shore of 1,700 ha Douglas Lake (Fig. 1). We present MAT as the average of daily maximum and minimum values and MAP as the sum of daily water-equivalent values (UMBS 2024). For atmospheric deposition, we present annual total inorganic N deposition values, as obtained from the National Atmospheric Deposition Program, of which UMBS is site MI09.

2.4 Data analysis- This paper is intended to provide a holistic view of interactions between ecosystem components over time, and how they influence C cycling. This required synthesizing and analyzing many different types of data according to a range of appropriate approaches and techniques. Analytical details are provided here, in Appendix S1, and in figure captions. The following narrative enumerates the specific statistical tests used to address each topic of interest; parenthetical references to figures are provided to aid in connecting these statistical tests to the results we present and the inferences we draw from them. To test for differences in total ecosystem C stocks (and in individual pools) over succession (Fig. 2a), we treated each plot in each chronosequence and old reference stand as an experimental unit, binned plots into 40-year age classes, and used one-way ANOVAs with Fisher's Least Significant Difference post-hoc tests to identify significant differences between age classes. Changes in composition over the same 40-year age classes (Fig. 2b) are expressed as the proportion of aboveground live tree biomass in each of three functional groups (short-lived deciduous, long-lived deciduous, long-

339 lived coniferous); these data are presented for visualization but are not analyzed statistically. To
 340 test relationships between canopy structure and C uptake over succession (Fig. 3), we treated
 341 each plot in each chronosequence and old reference stand as an experimental unit, selected NPP_w
 342 as the response variable, and used multiple linear regression to determine the predictive capacity
 343 (as partial r^2) of leaf area index (LAI), maximum canopy height, and rugosity. We conducted
 344 separate regressions for young (22-48 years; $n=11$), middle-aged (66-84 years; $n=6$), and old
 345 (>100 years; $n=12$) stands. We standardized the values for the three structural parameters by
 346 subtracting their means and dividing by their standard deviations. To examine successional
 347 trajectories of canopy production in taxonomic detail in recent decades (Fig. 4), we treated
 348 individual plots in the reference and treatment tower footprints as experimental units, selected
 349 leaf litterfall mass as the response variable, and used year as the predictor variable in simple
 350 linear regressions to predict temporal trajectories in leaf production (total across all species), and
 351 for individual species (red oak), genera (red +sugar maples), and functional groups (early-
 352 successional species, i.e., aspen and birch). To test for successional trends in soil respiration (Fig.
 353 5), we treated each chronosequence and old reference stand as an experimental unit, and used
 354 simple linear regression to test for temporal trends (i.e., as a function of stand age) in Rs fluxes
 355 and proportions. We ran separate regressions for annual vs. growing-season only trends, for each
 356 of the following fluxes and proportions: 1) total soil respiration [Rst], 2) autotrophic soil
 357 respiration [Rsa], 3) heterotrophic soil respiration [Rsh], and 4) heterotrophic proportion
 358 [Rsh/Rst]. To examine relationships between stand age, fine root production, and soil respiration
 359 (Fig. 6), we conducted two regressions: first, stand age vs. mean growing season fine root
 360 production (averaged to the stand level across all available years of ingrowth core incubation);
 361 second, mean growing season fine root production vs. annual total soil respiration. To examine

362 long-term successional patterns in O horizon C stocks on the Burn Plots (section 3.2 narrative),
 363 we used unweighted effect size meta-analysis to test (1) for a significant effect of past
 364 disturbance on C stocks using categorical meta-analysis and (2) whether O horizon C stock
 365 changes vary significantly with time since disturbance using continuous meta-analysis. To test
 366 for long-term changes in soil properties across the Burn Plots (Fig. 7a), we treated individual
 367 stands as experimental units, selected A horizon pH, %C, and microbial biomass C as response
 368 variables, and used stand age as the predictor variable in simple linear regressions. To test for
 369 relationships between soil properties and biogeochemical function (Figs. 7b, c), we treated
 370 individual Burn Plot stands as experimental units, selected α -glucosidase activity rate as the
 371 response variable, and regressed it against pH and %C as predictor variables (respectively) using
 372 simple linear regressions. To test for changes in fungal community composition over succession
 373 on the Burn Plots (Fig. 8), we treated individual soil samples from each stand as experimental
 374 units. In QIIME, we (1) identified fungal families driving community dissimilarity across stands
 375 using ANCOM, (2) visualized fungal communities with a PCoA biplot, and (3) tested for
 376 statistical significance of community dissimilarity along a Bray-Curtis dissimilarity matrix. To
 377 identify soil properties driving community dissimilarity (Fig. 9), we again selected individual
 378 Burn Plots soil samples as experimental units and used a (1) dbRDA biplot to visualize
 379 dissimilarity in soil properties across stand ages, and (2) forward stepwise regression, including
 380 α -glucosidase activity, microbial biomass C and N, soil $\text{NH}_4\text{-N}$ availability, soil C and N
 381 concentrations, pH, and water content as predictor variables, to identify statistically significant
 382 predictors of variation in community dissimilarity. To test for significant differences in
 383 competition for N among different EcM fungal families (Fig. 10), we treated individual plots in
 384 the reference and treatment tower footprints as experimental units, selected the ^{15}N value of bulk

fungal sporocarps from each family as the response variable, and used family as the predictor
 variable in a one-way ANOVA. To test for significant changes in mineral C stocks in recent
 decades (Fig. 11), we treated individual soil profiles as experimental units, selected total C stock
 as the response variable, and used a two-way ANOVA to test for significant effects of horizon
 (A, E, B), sampling year (1980, 2009), and their interaction on C stocks. To test for significant
 temporal trends in GPP, R_{eco} , NEE, and WUE (Figs. 12a-c), we used multiple linear regressions.
 For each response variable, we ran a multiple regression model with year (continuous) and
 treatment (dummy variable, coded relative to the reference footprint) to test for significant
 temporal trajectories and significant differences between footprints. Flux timeseries were 2000-
 2020 for the reference tower and 2008-2020 for the treatment tower. We used a paired t-test to
 determine whether mean cumulative growing season soil respiration efflux differed in reference
 vs. treatment tower footprints over the long-term timeseries. To test for significant temporal
 changes in MAT and MAP (Figs. 12d, 12e) we ran simple linear regressions with year as the
 predictor, for a 1980-2020 climatology period. To test for significant changes in atmospheric N
 deposition, we used piecewise regression to examine break-point timing and slope for annual
 total inorganic N deposition values from the UMBS NADP station, 1980-2020.

In the case of all tests, we used data transformations as necessary to mitigate non-normality or
 unequal variances, and set $P < 0.05$ as the threshold for accepting test results as statistically
 significant. In a limited number of cases and in the context of results significant at $P < 0.05$, we
 discuss several results with $P > 0.05$ but $P < 0.10$ as marginal trends. We completed t-tests,
 ANOVAs, and regressions in SigmaPlot 14.0 (SYSTAT Software, San Jose, CA US). We used
 MetaWin 3.0 (Rosenberg 2022) to analyze Burn Plots soil C data. Data processing and analyses
 of fungal community composition were completed in R and QIIME.

3. Results

3.1 Ecosystem C stocks and aboveground dynamics over succession- Total ecosystem C stocks in UMBS upland forests increase over 200 years of succession ($P < 0.001$), largely due to increases in aboveground biomass ($P < 0.001$), which exceed soil as the largest C pool by ~50 years of stand development (Fig. 2a). Smaller C pools also show significant temporal dynamics, including larger standing dead tree C stocks in the 40-80 and 80-120 year age classes ($P < 0.001$), larger O+A horizon C stocks in 120-160 and >160 year old stands, and larger DWD ($P < 0.001$) and smaller fine root ($P = 0.05$) C stocks in >160 year old stands (Appendix S1). As this successional increase in total ecosystem C unfolds, forest composition changes, with short-lived pioneer species giving way to longer-lived deciduous or coniferous taxa (Figure 2b).

Rates of biomass increase (as NPP_w) also vary over succession, with the aspects of canopy structure influencing NPP_w shifting over time (Fig. 3). Early in succession (plots in stands that are 22-48 years old), NPP_w varies nearly four-fold between plots, but this variation is unrelated to structural parameters (multiple regression $P > 0.05$). In middle-aged stands (66-84 years old) NPP_w is less variable among plots, and is significantly related (multiple regression $P = 0.03$) to canopy height (partial $r^2 = 0.72$), leaf area index (partial $r^2 = 0.21$), and, to a lesser extent, canopy rugosity (partial $r^2 = 0.04$). Old stands (>100 years old) possess the widest between-plot variance in NPP_w , which is significantly related (multiple regression $P = 0.02$) to leaf area index ($r^2 = 0.60$), and to a lesser extent canopy height and rugosity (both partial $r^2 < 0.05$).

Leaf litterfall data from the flux tower footprints detail the changes in forest composition that occur as pioneer species give way to longer-lived taxa at the close of the first century of succession. In the reference tower footprint where background natural succession is ongoing, a 23-year decline in leaf production by early-successional aspen and birch (Fig. 4A; regression

P<0.001) has been offset by a significant increase by red oak (Fig. 4B; P<0.001), maintaining whole-canopy leaf production at the same level (Fig. 4A; regression P>0.05). Leaf production by maple (primarily *A. rubrum*, secondarily *A. saccharum*) has remained unchanged over this interval (Fig. 4B; regression P>0.05). The 14-year litterfall record for the treatment tower footprint reveals a pattern of compositional change that is functionally similar, but greater in rate and magnitude. There, where the mortality of early-successional taxa was accelerated by stem girdling (Fig. 4A), leaf production by oak and maple both increased significantly (Fig. 4C; P<0.001), albeit threefold more rapidly for oak, based on slope coefficients for the two taxa. Indeed, oak transitioned from being subdominant to maple during the first 5 years of litterfall collections in the treatment footprint, to consistently being the canopy dominant over more recent years.

3.2 Belowground C, soil biogeochemistry, and microbial dynamics over succession- Soil respiration is the largest loss term in the UMBS C budget, and is dominated by emissions during the growing season (Fig. 5). On an annual basis, total soil respiration (Rst) does not vary with successional stage (P=0.29), but marginal decreases in autotrophic (Rsa; P=0.07) and increases in heterotrophic (Rsh; P=0.06) components lead to an overall increase in the proportion of total emissions that are heterotrophic (Figs. 5a, 5b). During the growing season, successional trends in soil respiration are more evident, with Rst (P=0.04) and Rsa (P=0.03) declining with stand age, and Rsh/Rst increasing with stand age (Figs. 5c, 5d). Growing season fine root production, which is unrelated to stand age (Fig. 6a, P=0.22), is in fact a stronger predictor of Rst than stand age itself (Fig. 6b; P<0.01).

Soils data from the Burn Plots chronosequence reveal changes in soil C, biogeochemical, and microbial properties over the course of succession. Categorical meta-analysis of the 5 known

sampling campaigns shows that O horizon C stocks were diminished by the clearcut+burn that established these stands (-29%, bootstrapped confidence interval -14% to -42%), with no temporal trend to imply recovery over the duration of the chronosequence (meta-regression, $P=0.74$). In contrast to O horizons, topsoils appear to be recovering as stands develop over the course of the chronosequence. Successional trends (Fig. 7a) include increased soil %C ($P=0.03$) and microbial biomass C ($P<0.01$), and decreased pH ($P=0.03$). These trends in turn have biogeochemical consequences, with higher %C and lower pH being linked to higher rates of cellulose-degrading enzyme activity (α -glucosidase; Figs. 7b, 7c; $P=0.03$).

Across the Burn Plots, successional changes in soil pH, %C, microbial biomass, and α -glucosidase activity are accompanied by significant changes in fungal community composition (Fig. 8; $P=0.025$). During the initial decades after disturbance, taxa in the EcM Amanitaceae and mixed saprotrophic/EcM Clavariaceae dominate. Stand development progresses through stages dominated by EcM taxa in the Tricholomataceae and Russulaceae, and by a century of succession, the fungal community is dominated by EcM taxa in the Cortinariaceae (Fig. 8; see Appendix S1 for specific taxa). This long-term fungal community shift has been accompanied by the topsoil property changes in Fig. 7, as well as increased moisture and root biomass, and a successional decline in soil $\text{NH}_4\text{-N}$ availability (Fig. 9; $P<0.05$). Of all these soil properties, pH, moisture, α -glucosidase activity, root biomass, and $\text{NH}_4\text{-N}$ availability are most strongly related to shifts in the fungal community over the course of succession on the Burn Plots (Fig. 9).

Patterns from the Burn Plots are informed by two results from other experimental designs that address fungal N cycling and soil C stocks. First, fungal sporocarp analyses from the decadal ^{15}N tracer experiment in the ~110 year old forests of the flux tower footprints reveal that of the four most frequently observed EcM families, sporocarps of taxa in the Cortinariaceae have

significantly higher ^{15}N enrichment (Fig. 10; $P=0.01$). Second, multidecadal soil monitoring in the 1911-origin reference stand adjacent to the Burn Plots demonstrates significant declines in A and E horizon C stocks in the last 30 years in this ~110 year old reference stand (Fig. 11).

3.3 Ecosystem C fluxes and climate forcings- Two decades of flux tower observations reveal significant changes in whole-ecosystem C fluxes and their environmental drivers (Fig. 12). Ecosystem C uptake (GPP, negative=uptake) has become significantly more negative with time, with no difference in the rate of change between reference and experimentally accelerated succession footprints (Fig. 12a, lower portion). Concurrently, R_{eco} has increased significantly in the reference footprint, but not in the treatment footprint (Fig. 12a, upper portion). In both footprints, WUE has increased concurrently with GPP, with no difference in the rate of increase between footprints (Fig. 12b). In terms of net C balance, NEE has become more negative with time in both footprints, but this increase in net C uptake has been significantly weaker in the background natural succession footprint due to the long-term increase in R_{eco} (Fig. 12c). Over the common period of record, cumulative mean growing season R_{st} has been significantly larger in the reference footprint than the treatment footprint ($P<0.001$, Appendix S1).

Changes in C and water fluxes observed in the past 15-20 years have been accompanied by ongoing changes in climate and atmospheric deposition over the past 40 years. From 1980 to 2020, UMBS became significantly warmer (Fig. 12d) and wetter (Fig. 12e), with increases of $+0.25^\circ$ MAT and $+5.0$ cm MAP per decade. Over the same period, wet inorganic N deposition declined significantly (-25% ; Fig. 12f), primarily since the year 2000.

4. Discussion

4.1 Synthesis of ecosystem changes over succession

Our synthesis illustration (Fig. 13) integrates this multitude of ecosystem components, their changes and interactions over the range of timescales and experimental designs at UMBS. This illustration is a simplified representation of results reported above, which are parenthetically referenced in the text below and complemented with UMBS-relevant citations.

Following stand-replacing disturbance on the sandy outwash plains of UMBS, aboveground biomass accumulation is initially rapid and unrelated to canopy structural development (Figs. 2a, 3b, 13a). As stands approach mid-succession, vertical canopy development drives the accumulation of aboveground biomass, which soon exceeds soil as the largest ecosystem C pool (Figs. 2a, 3b). Increases in aboveground biomass C continue later into succession, where between-stand variation in leaf area explains rates of continued increase (Figs. 2a, 3b, 13c).

Stand-replacing disturbance diminishes O horizon C stocks (section 3.2, meta-analysis) for 100 years, but soil C and related soil physicochemical, biogeochemical, and microbial properties begin to recover during the first century of succession. Recovery encompasses increases in A horizon moisture, exchangeable acidity, root and microbial biomass, and Bs horizon pedogenic iron concentrations, and decreases in A horizon pH, base cations, cation exchange capacity, and soil N availability (Figs. 7a, 9, 13a; Nave et al. 2019; White et al. 2004). These soil property changes have functional consequences including increased rates of α -glucosidase activity and a successional increase in the heterotrophic fraction of soil respiration (Figs. 7b, 7c, 5a, 5c, 13a), and are accompanied by long-term shifts in fungal and tree community composition (Figs. 13a, 13b). Over this 100-year timescale, communities shift from being dominated by short-lived aspen and birch with mixed saprotrophic/enzymatically weak EcM fungal communities, to increased representation of longer-lived deciduous trees and fungal communities dominated by EcM taxa in the Cortinariaceae, with known organic matter decomposition abilities (Figs. 2b, 8,

13a, 13b; Bödeker et al. 2014; Kyaschenko et al. 2017). As these compositional shifts unfold over decades, the heterotrophic fraction of total soil respiration increases (Figs. 5a, 5c). After a century of succession, combined O + A horizon C stocks recover from stand-replacing disturbance (Fig. 13c).

By 100 years of succession after stand-replacing disturbance, aspen and birch have begun to die out, at rates that naturally vary according to soil and landform properties, and in the case of the experimental girdling disturbance, have been deliberately accelerated (Figs. 2b, 4a, 13b; Nave et al. 2014). Similar to the Burn Plots but over a shorter timescale, accelerated aspen-birch mortality in the treatment flux tower footprint causes a temporary increase in soil N and an accompanying excursion in fungal community composition, with increased saprotrophic and decreased EcM taxa (Fig. 13b; Castillo et al. 2018; Nave et al. 2011a; 2013; 2014). As aspens die, oak outcompetes maple for canopy gaps and N, especially in the treatment tower footprint, with its EcM associates in the Cortinariaceae proving highly competitive for N (Figs. 4a, 4b, 10, 13ab; Nave et al. 2013). In reference forests where gradual background mortality of aspen and birch is ongoing (i.e., the reference flux tower footprint and the 1911-origin reference stand adjacent to the Burn Plots), soil and total ecosystem respiration increase as surface soil C stocks decline, resulting in a weaker overall C sink than in the experimentally accelerated succession footprint (Figs. 11, 12a top, 13b). In both footprints, GPP and WUE exhibit ongoing increases with mean annual temperature and precipitation (Figs. 12a bottom, 12b, 12d, 12e).

Predictions of C cycling and its drivers through the end of a second century of succession are hindered by the prospects of continued climate change and the unique conditions on which the three old reference forests in our experimental design are found (Fig. 13c). While some of the more robust chronosequence trends may hold into the future, such as long-term aboveground

biomass increases, (Fig. 2a), others are less clear. For example, differences in LAI and NPP_w between the ~130 year old pine-dominated stands on outwash and the ~190 year old hemlock-hardwoods on moraine may be successional trends, or results of unique landform and soil properties (Fig. 13c, Albert and Minc 1987; Nave et al. 2017; 2019; Scheuermann et al. 2018).

4.2 Detailed discussion of ecosystem components

4.2.1 Aboveground wood production and canopy structure- Our finding that the canopy metrics that drive aboveground growth change over the course of succession (Fig. 2) addresses the first of the predictions enumerated at the close of the Introduction. In young stands, canopy structural complexity does increase with age, but it is not the driving force for aboveground growth. Instead, the severity of the stand-establishing disturbance may be a more important determinant of between-stand variation in NPP_w (Scheuermann et al. 2018). Canopy structural control of NPP_w becomes stronger in middle aged stands, evidenced by low residual variance in the structure – NPP_w relationship, and suggests that the rate at which a deeper canopy develops fundamentally constrains NPP_w during this period of rapid vertical growth, possibly involving a tradeoff between vertical growth and lateral crown expansion (i.e., increased LAI). Beyond a century of development, variation in LAI explains the majority of the variance in NPP_w . This may indicate that as rates of overstory tree vertical growth begin to slow with succession, canopy stratification (including the developing understory) becomes more important to whole-canopy photosynthetic C fixation (Murphy et al. 2023).

4.2.2 Soil C – microbial interactions over succession- Soil C stocks and microbial communities were profoundly altered by region-wide stand-replacing disturbances a century ago, but have begun to recover during the century of succession that has unfolded since. The meta-analytic reduction in O horizon C stocks that we report (-29%, section 3.2) falls in the range of losses

reported in meta-analyses of harvest and fire for temperate forest soils generally (Nave et al. 2010; 2011b), but is larger than typical in the Lake States (Nave et al. 2021). This may reflect a compound effect of the second (experimental) cut-and-burn disturbance applied to create each Burn Plot (i.e., chronosequence stand) throughout the 20th century, which is in addition to the landscape-wide cutting and fires that occurred in the latter 19th through early 20th centuries (Gough et al. 2007). In contrast to persistently diminished O horizons, A horizons across the Burn Plots appear to be recovering, with %C, microbial biomass C, and pH showing significant relationships with stand age (Fig. 7a). These results independently confirm a previously reported decrease in topsoil pH across the Burn Plots (Nave et al. 2019), and add mechanistic insight by revealing that as topsoils begin their post-disturbance recovery of acidity and organic matter, the microbial community and its enzymatic functions (Fig. 7b, 7c) follow suit.

4.2.3 Coupled tree-fungal compositional-functional changes- Our synthesis of results from the Burn Plots and FASET reveals that stand-replacing and partial disturbances induce microbial and N cycling responses that are similar in pattern, but different in timescale and magnitude. On the Burn Plots, clearcutting + fire causes soil N availability and nitrophilic fungi (EcM and saprotrophs) to increase abruptly, then decline over 100 years as N-foraging EcM decomposers in the Cortinariaceae rise in dominance (Figs. 8, 9; Birkebak et al. 2013; Bödeker et al. 2014; Chen et al. 2019; Kyaschenko et al. 2017; Nave et al. 2019; White et al. 2004). Similarly, in FASET, the pulse of aspen and birch mortality in the treatment footprint led to a 2-4 year increase in soil N availability relative to the reference footprint, accompanied by a decline in EcM and an increase in saprotrophic fungi (Castillo et al. 2018; Nave et al. 2011a; 2013; 2014). Since that time, EcM fungi and their N uptake have recovered, with taxa in the Cortinariaceae outcompeting taxa in the Amanitaceae, Boletaceae, and Russulaceae for the ecosystem ¹⁵N tracer

that was applied during the disturbance period (Fig. 10). Based on their strong decomposing enzyme production, the shift in the fungal community towards the Cortinariaceae would seem to favor increased rates of soil C mineralization and CO₂ emission.

Successional changes in microbial communities and their functions go hand-in-hand with changes in tree community composition and functioning, addressing our second prediction (coupled aboveground-belowground compositional-functional linkages). Increases in the enzymatically proficient Cortinariaceae may mean that these EcM taxa are increasing the ability of longer-lived hosts (primarily red oak) to access less available forms of soil N (Jorgensen et al. 2022; Leduc et al. 2013; Lindahl et al. 2021). Indeed, red oak has outcompeted red maple for soil N during the successional transition out of aspen-birch dominance (Nave et al. 2013), has higher foliar N concentrations (Nave et al. 2009), rates of stomatal conductance, photosynthesis, and sap flux (Gough et al. 2021; Matheny et al. 2017). These acquisitive vs. conservative traits may underpin the multi-decadal trends in canopy production by the two species. Over this period, oak has had the clear competitive advantage (Fig. 4), especially in the treatment footprint, where water and N availability were elevated for 2-4 years following experimentally accelerated aspen mortality (He et al. 2014; Nave et al. 2014). Compared to the aspen they are replacing, however, oak and maple both have lower rates of photosynthesis and stomatal conductance, and lower foliar N concentrations (Gough et al. 2021; Nave et al. 2009). Thus, this successional transition to longer-lived species may have longer-term consequences for functions such as water use, C uptake and respiratory emissions at the ecosystem scale.

4.2.4 Ecosystem C fluxes, net balance, and environmental drivers- Over the course of succession, compositional change from aspen to longer-lived tree taxa with lower photosynthetic rates would seem to favor reduced whole-canopy GPP. Decades of monitoring in the flux tower footprints

shows that the opposite has occurred (Fig. 12). This addresses our third prediction and implies that external forcings, such as warming and wetting, are overriding compositional influences on successional trajectories of C cycling, at least on the uptake side. On the emissions side, the picture is more complex. Respiratory emissions should be decreasing with succession in the tower footprints, because the longer-lived taxa replacing aspen and birch have higher C use efficiency (less C respired per unit C fixed; Gough et al. 2021), and the chronosequences suggest long-term declines in R_{st} and R_{sa} with age (Fig. 5). Counter to that prediction, R_{eco} has remained stable in the treatment footprint where composition has changed rapidly over the last decade, and increased in the reference footprint. Some portion of this increase in R_{eco} in the reference footprint has been due to increased R_{st} emissions, which have been larger than in the treatment footprint in the last 1-2 decades (section 3.3. and Appendix S1). These elevated respiratory losses are consistent with surface soil C stock declines observed over recent decades in the 110 year old Burn Plot reference stand, which is in all regards similar to the UMB reference footprint (Fig. 11). Collectively, these patterns imply an overarching, climate-driven increase in respiratory emissions, which is being canceled out in the treatment footprint by rapid successional change to tree taxa with lower specific respiration rates.

Though it is unclear whether climate or composition has the stronger influence on its constituent fluxes, the increasing C sink strength (NEE) over time in both footprints indicates that GPP has increased more rapidly than R_{eco} (Fig. 12). This strengthening C sink over the past 1-2 decades of succession has been concurrent with increased WUE in both footprints, due in part to the replacement of aspen and birch with oak and maple, which have higher WUE (Matheny et al. 2014). However, similar trends in NEE and WUE are occurring in other humid north temperate forests where compositional change is less likely to be a substantial mechanism, implying a role

for broader environmental drivers such as warming, wetting, CO₂ fertilization, and N oligotrophication (Finzi et al. 2020; Groffman et al. 2018; Hollinger et al. 2021; Keenan et al. 2013). The key takeaway from this trend is thus much the same as for respiratory emissions: there is evidence for a broader climatic influence on ecosystem C balance and water use, and an influence of successional change in composition. Unfortunately, our experimental design cannot be parse these factors as independent drivers.

One factor that may be influencing respiratory emissions at UMBS is non-native earthworms. Most of the earthworm taxa present at UMBS in recent decades have been present since the early- to mid-1900s, but changes in their spatial distribution and community composition have occurred (Crumsey et al. 2014). One of the more notable changes has been the expansion of *Lumbricus terrestris*, which directly consumes and vertically redistributes larger quantities of surface leaf litter than other taxa (Crumsey et al. 2015). On balance, *L. terrestris* and other taxa, including those that consume and redistribute mineral soil organic matter, have a small negative effect on soil C UMBS (Crumsey et al. 2013). This implies that declining soil C stocks and increasing soil and ecosystem respiration is due, in some part, to the activity of these organisms. Overall, earthworm spatial distribution and community structural will likely continue to change over time at UMBS, in a patch dynamic influenced by factors such as road proximity, soil texture, and tree community composition (Crumsey et al. 2014). As these changes unfold, earthworm dietary preferences may interact with forest successional dynamics to affect C cycling outcomes, as maple leaf litter is much preferred, and more quickly consumed, than oak litter (Crumsey et al. 2013).

4.3 Context and broader relevance to north temperate forest ecosystems and their management-

Our study is both representative and unique within the context of its region, in ways that help to

understand its relevance to C cycling in other north temperate forests. In its composition, UMBS is broadly representative, as the aspen-birch cover type is the most extensive at the site and across the broader Laurentian Mixed Forest Province (Nave et al. 2017a; Ruefenacht et al. 2008). In its age, UMBS is unique, as aspen-birch forests tend to be younger because their persistence depends on large-scale disturbance or management that has not occurred at UMBS (Carson et al. 2023; Friedman and Reich 2005). In terms of broader relevance, many of the general patterns that we report here are similar to other empirical studies across the U.S. Lake States, Northeast, and adjacent Canada. These include a strengthening C sink in a maturing forest over recent decades (Finzi et al. 2020; Hollinger et al. 2021), successional increases in ecosystem C stocks that are driven by aboveground biomass (Alban and Perala 1992; Keeton et al. 2011), and long-term recovery of soil properties following disturbance (Poirier et al. 2016; Roy et al. 2021). However, for every one of these general patterns across the region, there are studies that show divergent or more nuanced trends (Arain et al. 2022; Desai et al. 2023; Gao et al. 2018; Prest et al. 2014; Wang et al. 2014). This highlights that even if some successional trends are predictable (e.g., increases in total ecosystem C) and some drivers are known (e.g., leaf area, soil texture), site specifics (e.g., disturbance history, management prescriptions) can still be equally or more important (Ford and Keeton 2017; Hoover 2011; Kern et al. 2021; Latty et al. 2004).

In the north temperate forest literature, the lack of any one consistent successional C trajectory or set of drivers reflects diversity in forest ecosystems, including in the ways that they have been managed or otherwise altered by people. Accepting this diversity and the uncertainty that it introduces into future forest functioning is part of forest management. Of particular note in this context are the many factors we show other than stand age that influence C cycling, and the possibility that these factors can serve as explicit targets for manipulation when C is a

management objective. Managing forests means much more than managing their age, e.g., by clearcutting to restart, or allowing to “mature” by leaving them untouched. Directly and indirectly, managing forests means manipulating structure (above and belowground), composition (plants and microbes), and relationships between ecosystem components, including their functional and biogeochemical outcomes. Taking this ecological view of forest management, and setting aside the notion that forest age is a driver itself, we offer several management applications relevant to north temperate forests.

The most timely management application of our results emerges from the decadal trends in C uptake and emissions in the two flux tower footprints. In the treatment footprint, the intentional killing of over one-third of the live trees has had a paradoxical effect: an increase in C sequestration. There, altering forest composition redistributed resources (access to light, soil N) from resource-demanding to resource-efficient taxa. This action decreased canopy structural complexity—a driver usually positively associated with C sequestration (Gough et al. 2021; Hardiman et al. 2011; Thom and Keeton 2020)—and nonetheless increased C sequestration, implying a greater role for N than light limitation of C uptake. In contrast, in the reference footprint, natural succession towards an older and more structurally complex condition has been accompanied by a weaker increase in C sink strength, due to a concurrent increase in respiratory losses. Collectively, this difference between footprints implies that partial harvests to manipulate composition may be a way to simultaneously increase within-forest C sequestration, while removing C that would otherwise be respired and instead storing it in harvested wood products (Gahagan et al. 2015; Powers et al. 2011). Our results suggest such treatments will be most effective where there is a wide difference in resource efficiency between the dominant trees that are removed, and the codominant or suppressed trees that are released, with greater trait diversity

in released species facilitating climate adaptation and C mitigation (Clark and D’Amato 2021; Thom et al. 2021; Wiechmann et al. 2022).

Our findings provide a belowground view on the typically aboveground-focused work of managing forests. We show that predictable tree community composition changes over succession are linked to concurrent changes in soil properties and microbial communities. Whether severe and stand-replacing or partial and patchy, disturbances that kill trees accelerate the N cycle and shift the fungal community from low-N EcM fungi that are strong decomposers, to more nitrophilic EcM and saprotrophic fungi. These shifts can be subtle and short-lived, as in the experimental girdling that accelerated succession in FASET, or more persistent, as in the clearcut harvest + fire of the Burn Plots, where fundamental soil properties are still changing many decades later. In both cases, aboveground dynamics (tree growth, community composition) have successional implications for belowground processes, including root production, soil CO₂ emissions, and long-term soil C stock changes.

The final management consideration of our synthesis emerges from its landscape context. At 4,000 ha, UMBS is large enough to host multiple experimental designs on the same kind of “site”—sandy outwash plains—yet also large enough that its landscape holds other ecosystem types where findings from these study sites do not apply. Thus across UMBS and other humid temperate glaciated landscapes with mosaics of mixed hardwood-conifer forests, our results will be most relevant on drier landforms and soils with lower nutrient availability. On more productive landforms and soils, successional patterns and C cycle trajectories may be different, or similar but for different underlying reasons. Drawing management applications from this and other long-term studies will therefore be most effective when disturbances and succession are seen as whole-ecosystem processes, with bottom-up and belowground factors given equal weight

to the plant communities that usually capture the most attention (Barnes et al. 1982). Vegetation treatments to achieve C cycle objectives can then be optimized to the kinds of ecosystems where they are most likely to be effective. For example, our results imply that partial harvests to release codominant or overtopped oak can increase ecosystem C sequestration, but that this outcome is most likely on drought-prone, low-nutrient soils where its water- and N-acquisitive traits make it best suited to capitalizing on release from competition. On more mesic, productive soils, treatments to release maple and capitalize on its high levels of water and C use efficiency may be more effective at increasing C sequestration. Still other management strategies, approaches, and tactics may be needed to balance C with other objectives in other settings. In this way, moving beyond stand-level thinking to landscape management mosaics can diversify the composition, structure, connectivity, climate adaptation capacity, and C management strategies of larger forested areas (Ontl et al. 2020; Thom and Keeton 2020). Ultimately, as climate, disturbance regimes, and species distributions continue to change, management must be conducted in this larger context in order to maintain forest functions- and in some areas, forests at all.

4.4 Key unknowns- Our work struggles with several limitations, and also highlights several areas in need of future research. Its most significant limitation emerges from its greatest strength, in the sense that so many experimental designs and datasets can be brought to bear on so many aspects of the ecosystem at a single site. While this allows for a holistic view of C cycling and its drivers, the lack of consistency in individual studies precludes an elegant, balanced synthetic analysis. The resulting narrative synthesis is as complex as the ecosystem itself, and in simplifying this complexity for clearer communication, we have minimized attention to potential issues such as stand-level variation in disturbance histories and the disconnect between inferences from a 20-year flux record vs. a 200-year chronosequence. The spatial and temporal

distribution of past clearcutting and fire at UMBS is more nuanced than we communicate here, and in focusing on what is consistent across disturbance types, we overlook the potential that localized past events may be influencing the results we report (Cooper 1981; D'Amato et al. 2017; Farmer 1958; Kilburn 1957). Similarly, we emphasize consistencies, rather than differences between the decadal patterns of succession in the flux tower footprints and the centuries-long continuum of chronosequence and old reference stands. Notably, our NEP estimates over these longer timescales indicate that only one of these stands is definitively a C sink (Clay et al. 2022), a result which is at odds with tower-based observations showing a consistent C sink over time.

One of the key unknowns in this study is the degree to which several unaddressed belowground processes and C fluxes vary over successional time. In this work, we are limited to relatively coarse metrics of belowground C cycling, such as soil respiration and fine root production. More detailed investigations of processes such as root exudation, fungal hyphal production, and bacterially-mediated C and nutrient transformations would result in more mechanistic inferences into microbial and biogeochemical drivers of belowground C cycling over succession. This is particularly true for N cycling processes, given the likelihood that this site's strong N limitation will only increase with CO₂ fertilization and N oligotrophication (Groffman et al. 2018).

Another key unknown in our study is the broader spatial (and longer-term temporal) relevance of our results pertaining to the ecophysiology and growth trajectories of red oak. The similarity of oak's increase at UMBS and the long-term Harvard Forest AmeriFlux site (Finzi et al. 2020), concurrent with the increasing C sink strength at both sites, implies a potential connection between composition and C cycling functions, not to mention a bright future for oak. In contrast, a recent synthesis of oaks in the context of climate change indicates more complex, and likely

negative prospects for the genus. Namely, the same acquisitive traits that have facilitated oak's competitive rise at these humid sites predispose them to decline as warming continues and vapor pressure deficit increases (Novick et al. 2022). Here, there are clear management implications for how drought, pests and pathogens, and C cycling interact in oak forests throughout the East. As climate change and other pressures continue to increase, management interventions such as thinning and prescribed fire may be necessary to support oak persistence, while accepting novel structures, reduced densities, and increased mortality as inevitable (Clark et al. 2022, Isaacson et al. 2023; Refsland et al. 2020). As conditions continue to change, further research will be needed to examine the ecological mechanisms by which management affects composition and C cycling, across a range of structures and successional stages, in oak and other forest types.

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Figure Captions

Figure 1. The UMBS landscape. Map shows the locations of plots within the paired chronosequence stands (1x cut+burn and 2x cut+burn) and old reference forests, as well as the US-UMB (reference) and US-UMd (treatment) flux tower plot networks. Shading indicates landscape-level physiography. Inset: UMBS' regional location.

Figure 2. Panel A: Total ecosystem C stocks and their distribution among pools, for plots in both chronosequences and the three old reference forests. Individual plots have been binned into 40-year age classes. Bars are means, errors are standard deviations, and letters denote significant differences between means (Fisher's L.S.D.; overall ANOVA $P < 0.001$). See Appendix S1 for statistical test results. **Panel B:** Proportions of total aboveground live tree biomass for three functional tree groups, by 40-year age class (presented for qualitative patterns but not statistically analyzed).

Figure 3. Panel A: NPP_w (mean and standard error) vs. stand age for young (22-48 years), middle-aged (66-84 years), and old (>100 years) stands. **Panel B:** For a multiple regression model for each age category, the proportion of variance in NPP_w explained by individual structural parameters, including leaf area index (LAI; blue), mean maximum canopy height (Max Height; orange), and canopy rugosity as variance of combined vertical and horizontal leaf distributions throughout the canopy (Rugosity; green). The residual (unexplained) variance in each regression model (young, mid, old) is plotted in gray. **Panel C:** Standardized structural parameter values and NPP_w rates for individual plots in young, middle-aged, and old stands. Structural parameters significantly related to NPP_w are indicated with enlarged points and best-fit lines.

Figure 4. Long-term annual leaf litter production in reference and treatment (accelerated succession) tower footprints. Leaf litter production is shown for all vs. only early-successional species (panel A), and for *Q. rubra* vs. *Acer* spp. (panels B, C). Points plotted are means and standard errors for litterfall collections in 10-14 plots per footprint. Best fit lines indicate statistically significant relationships between year and leaf litterfall flux. Note differences in the period of record for reference (1997-2019) vs. treatment (2006-2019) footprints.

Figure 5. Soil CO₂ emissions as a function of stand age. **Panel A:** Annual total soil respiration (RstA) flux and the proportion of the total from heterotrophic sources (Rsh/Rst). **Panel B:** Annual autotrophic (RsaA) and heterotrophic (RshA) soil respiration fluxes. **Panel C:** Growing season total soil respiration (RstGS) flux and the proportion of the total from heterotrophic sources. **Panel D:** Growing season autotrophic (RsaGS) and heterotrophic (RshGS) soil respiration fluxes. Values presented are stand-level means across the 2014-2021 growing seasons. Statistics are for simple linear regression, with best-fit lines added for statistically significant relationships with stand age. In panels A and C, filled circles are Rst fluxes and open circles are Rsh/Rst proportions. In panels B and D, green triangles are Rsa fluxes and purple squares are Rsh fluxes.

Figure 6. Panel A: Growing season fine root production as a function of stand age. **Panel B:** Growing season fine root production vs. annual total soil respiration. Points plotted are stand-level means across all available years of observation; P and r² values are for simple linear regression.

Figure 7. Panel A: Relationships between stand age and A horizon properties including total C (%; filled triangles) and microbial biomass C (mg kg⁻¹, open triangles) concentrations, and pH (filled circles). **Panel B:** relationship between A horizon %C and pH. **Panel C:** Relationship

1230 between A horizon pH and α -glucosidase activity. In each panel, r^2 and P values are presented
 1231 for simple linear regressions relating the individual variables.

1232 **Figure 8.** Principal coordinates analysis biplot of fungal community dissimilarity across A
 1233 horizon soil samples from the Burn Plots. Points and ellipses are color-coded according to stand
 1234 ages (in years, as of 2015 when samples were collected). Ellipses represent 95% confidence
 1235 intervals for each stand, and vectors represent the top 5 families responsible for community
 1236 dissimilarity.

1237 **Figure 9.** Distance based redundancy analysis biplot of fungal community and soil property
 1238 dissimilarity across A horizon soil samples from the Burn Plots. Points and ellipses are color-
 1239 coded according to stand ages (in years, as of 2015 when samples were collected). Ellipses
 1240 represent 95% confidence intervals for each stand, and vectors represent the soil properties that
 1241 were significantly related to community dissimilarity in a forward stepwise regression model.

1242 **Figure 10.** ^{15}N signatures of sporocarps collected from ^{15}N -labeled plots in reference and
 1243 treatment footprints. Plots were labeled with $^{15}\text{NH}_4\text{Cl}$ in 2010; sporocarps were collected in
 1244 2022. Plotted are means and standard errors for 5-15 sporocarps per family, with significant
 1245 differences ($P < 0.05$; Fisher's LSD) between families indicated with letters.

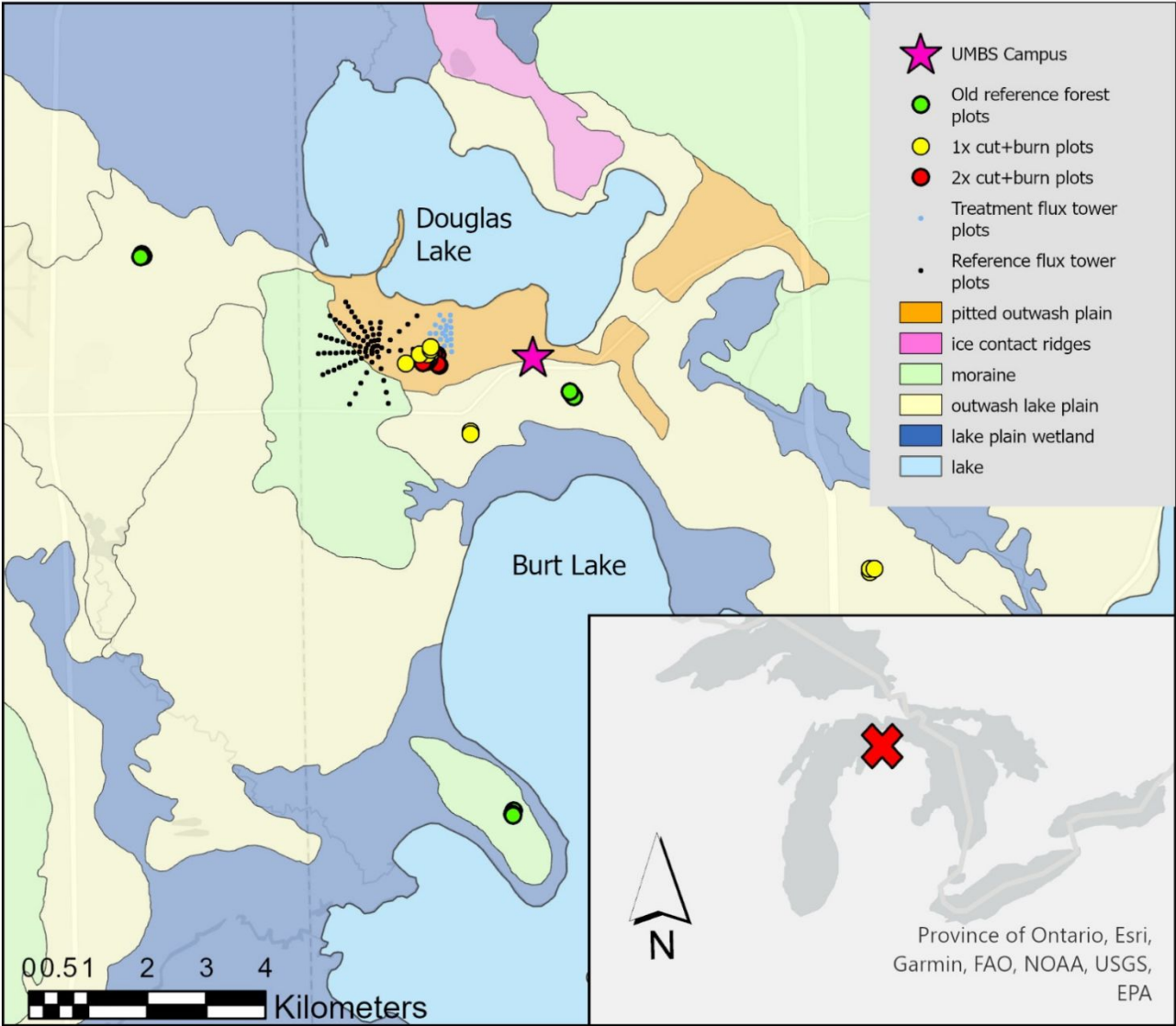
1246 **Figure 11.** Mineral soil carbon stocks, by genetic horizon, for soil profiles sampled in 1980 and
 1247 2009 in an area of unmanipulated 1911-origin reference forest adjacent to the Burn Plots. Plots
 1248 show means and standard errors; difference of means is significant ($P < 0.05$) for A and E
 1249 horizons.

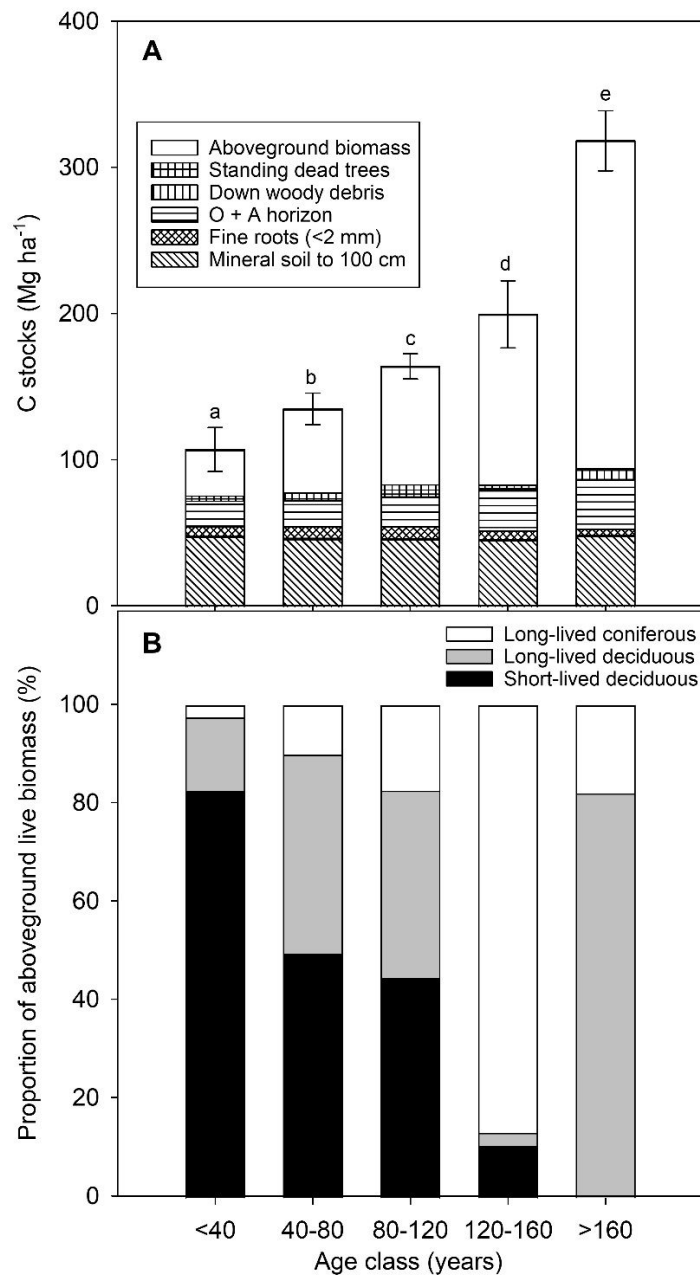
1250 **Figure 12. Panels A-C:** Ecosystem C fluxes and water use efficiency, for reference (filled
 1251 circles) and treatment (open circles) footprints. C fluxes follow eddy-covariance sign

conventions, such that uptake is negative (GPP, NEE) and emissions are positive (R_{eco}). To aid in visualization, timeseries lines connect individual annual values in each plot. Significance of year and treatment are denoted in each panel, with r^2 values provided when only year is significant, and the adjusted R^2 provided when year and treatment are both significant. **Panels D-F:** Mean annual temperature (D), precipitation (E), and bulk inorganic N deposition (F) at UMBS, 1980-2020, with regression P and r^2 values. Climatologic variables are averaged from daily observations from a weather station on the main campus, adjacent to Douglas Lake. Atmospheric deposition data are from the National Atmospheric Deposition Program (UMBS site ID MI09).

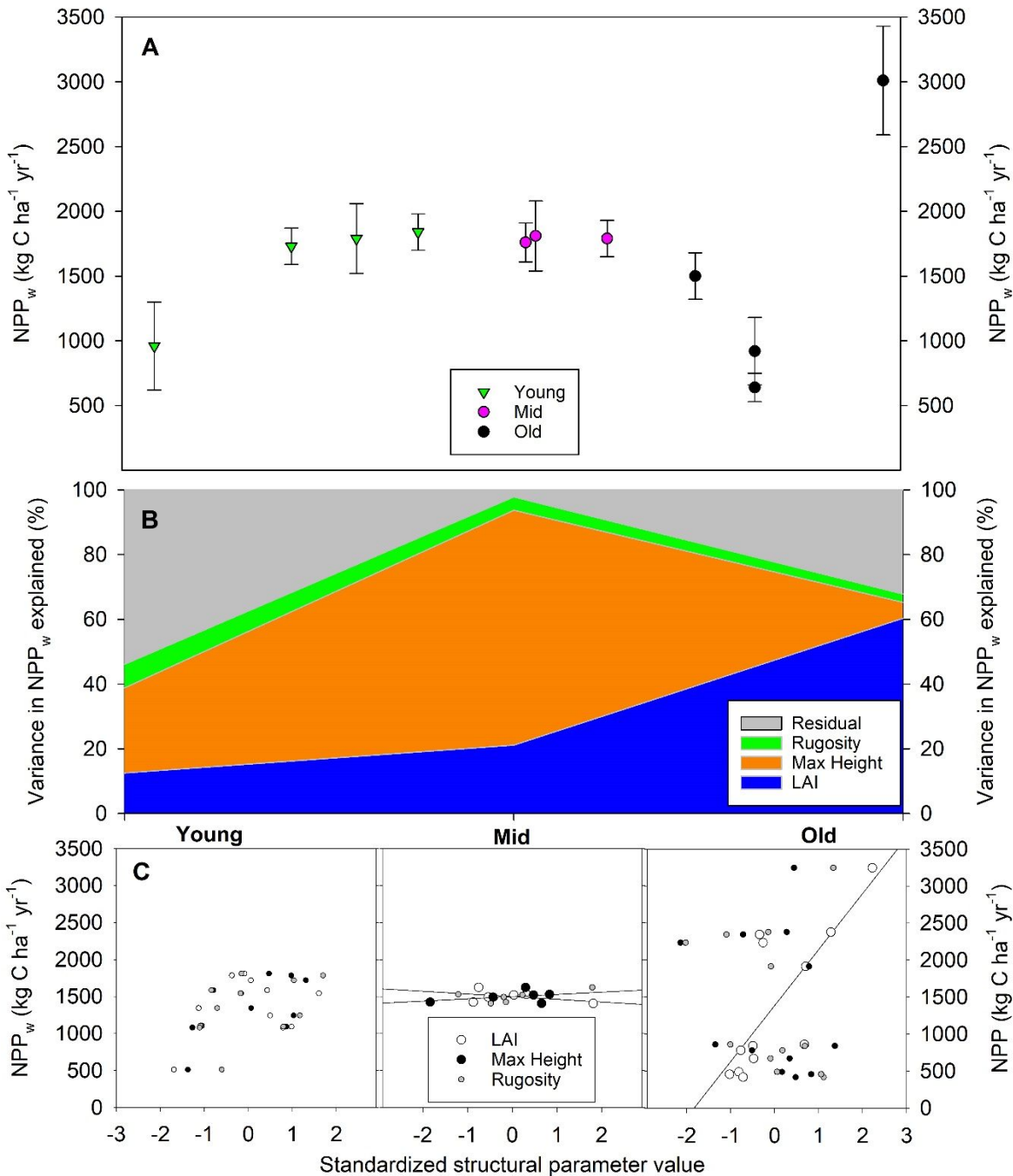
Figure 13. Synthesis of ecosystem changes over successional time. The three columns (A, B, C) correspond to the three experimental designs described in section 2.2, which correspond to three semi-discrete developmental phases that collectively span 200 years of succession. Column A shows trends inferred through the paired chronosequences (particularly the Burn Plots) over the first century of stand development. Column B corresponds to FASET, the paired comparison of background reference vs. experimentally accelerated succession footprints, over a 2-decade period from stand age ~90-100. Column C highlights the three old reference forests, with ages ranging from 130 years for the two pine-dominated stands to approximately 200 years in the uneven-aged hemlock-northern hardwoods stand. See section 4.1 for accompanying narrative.

1274 **Figure 1**



1281 **Figure 2**

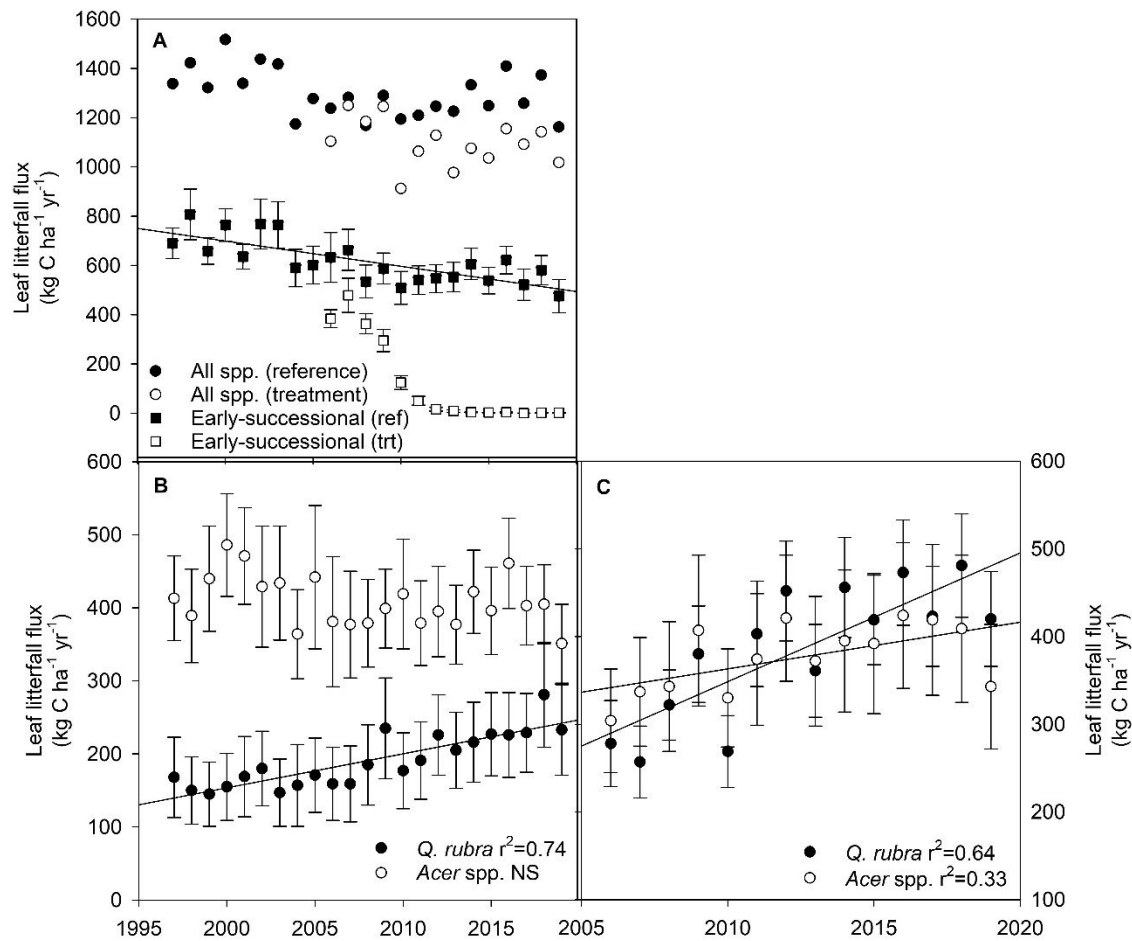
1286 **Figure 3**



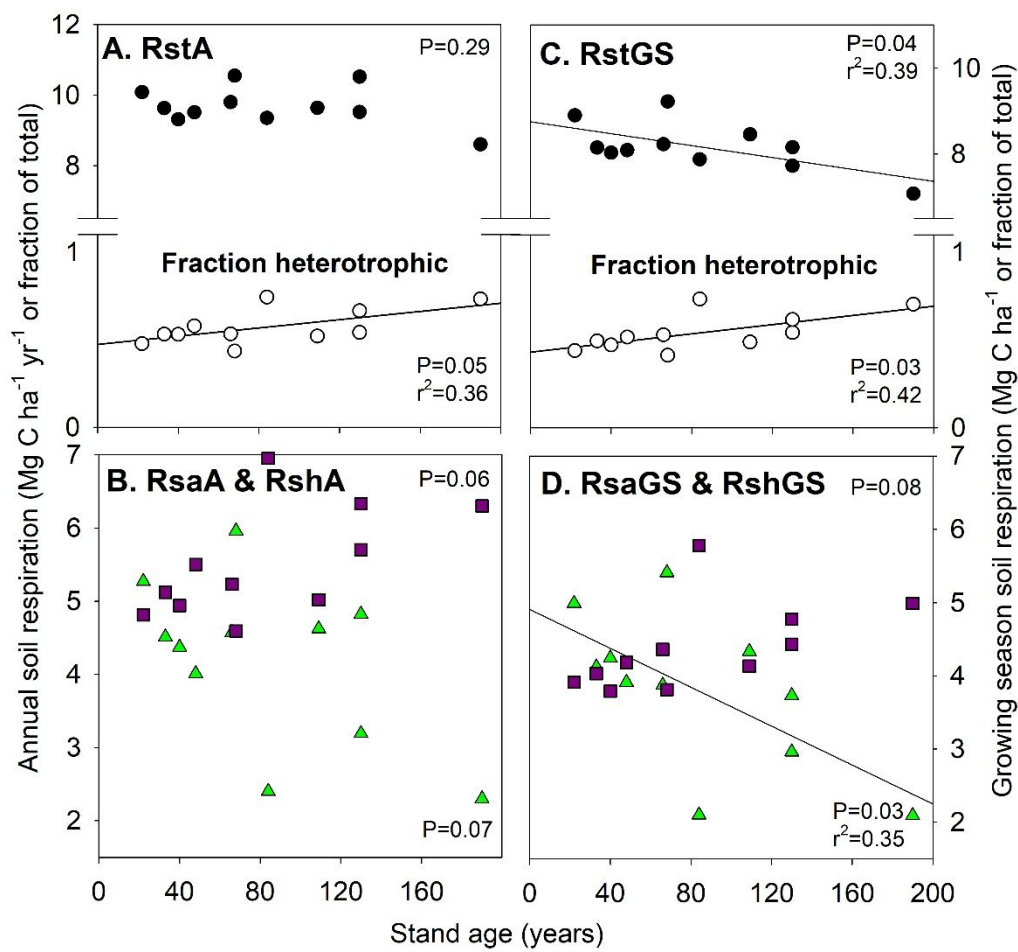
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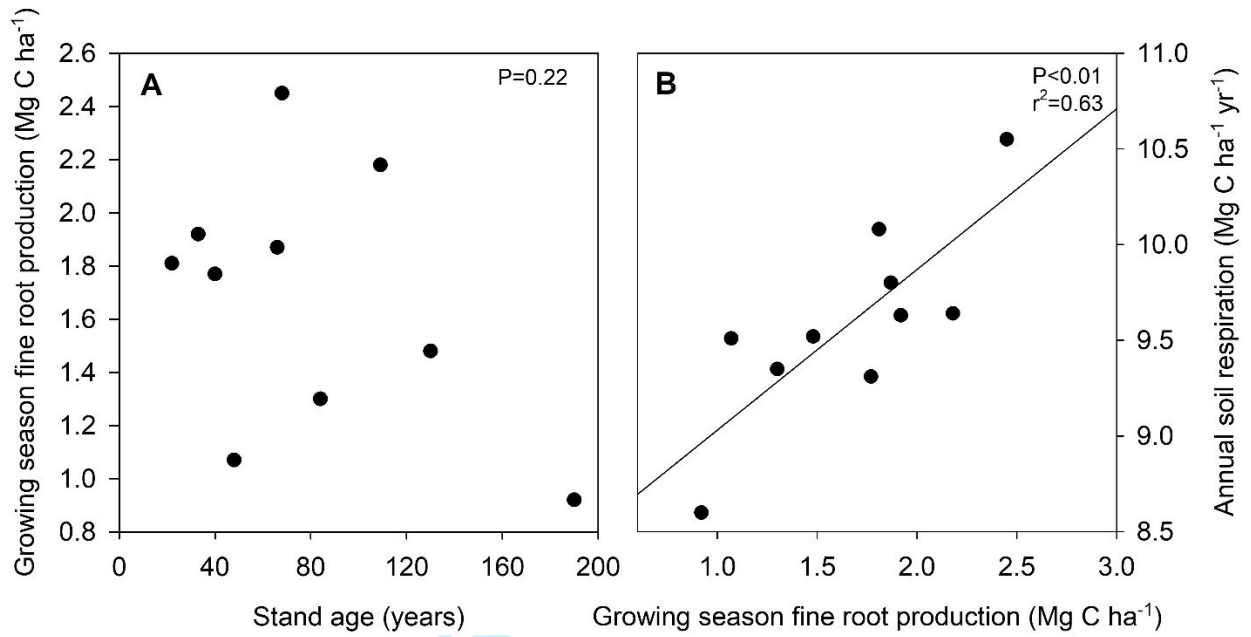
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Figure 4

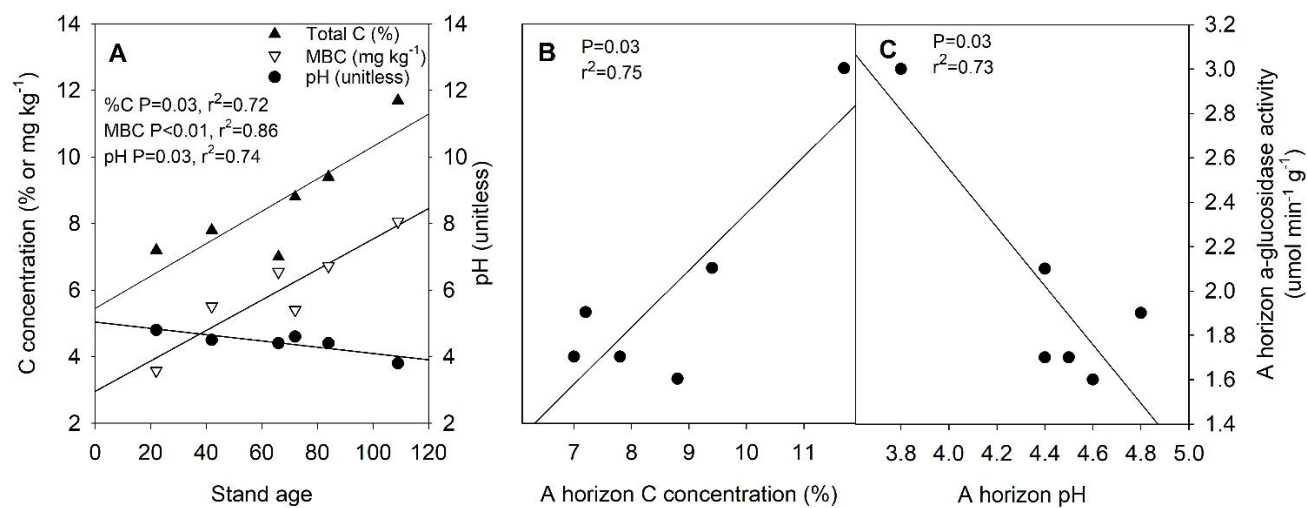
1299 **Figure 5**

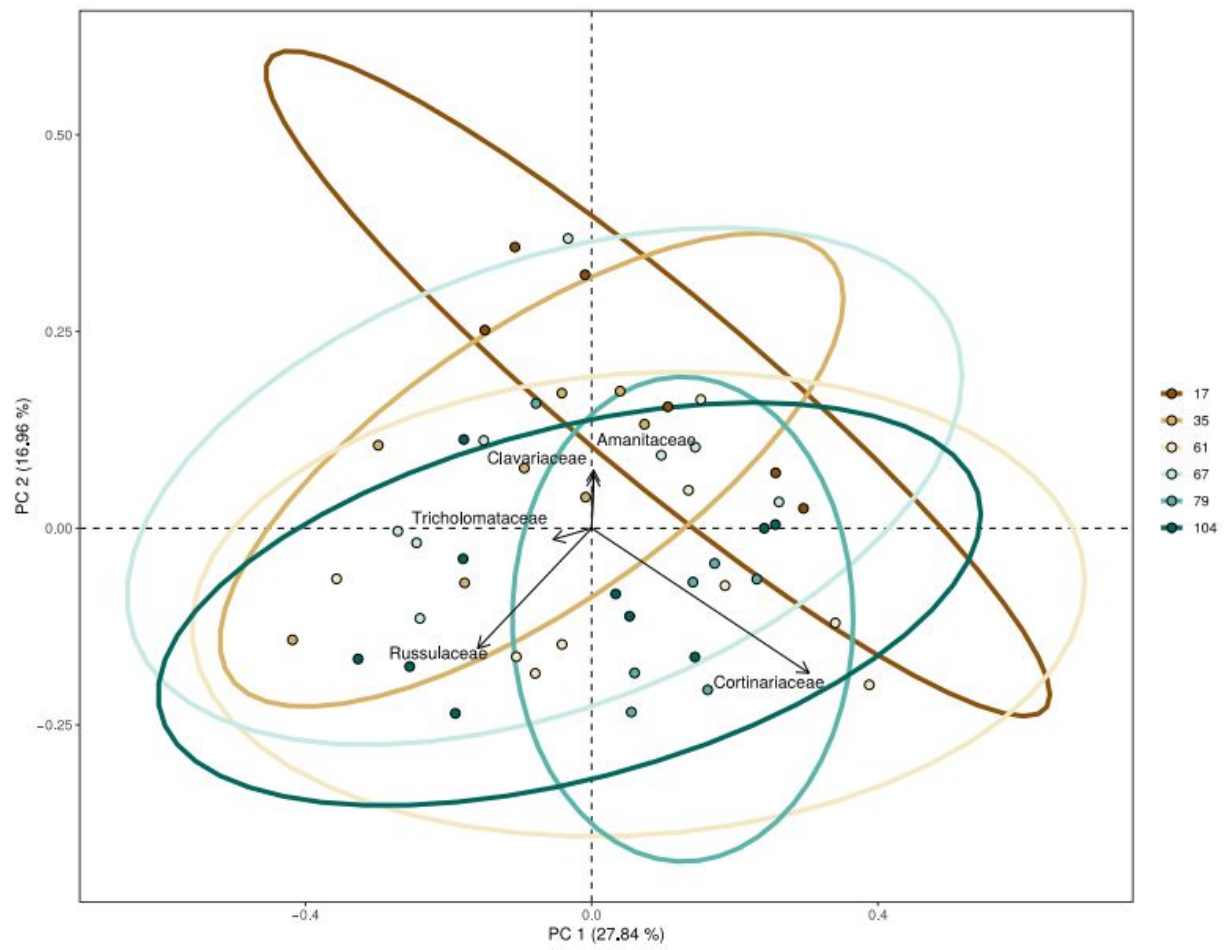


1307 **Figure 6**



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1332 **Figure 8**

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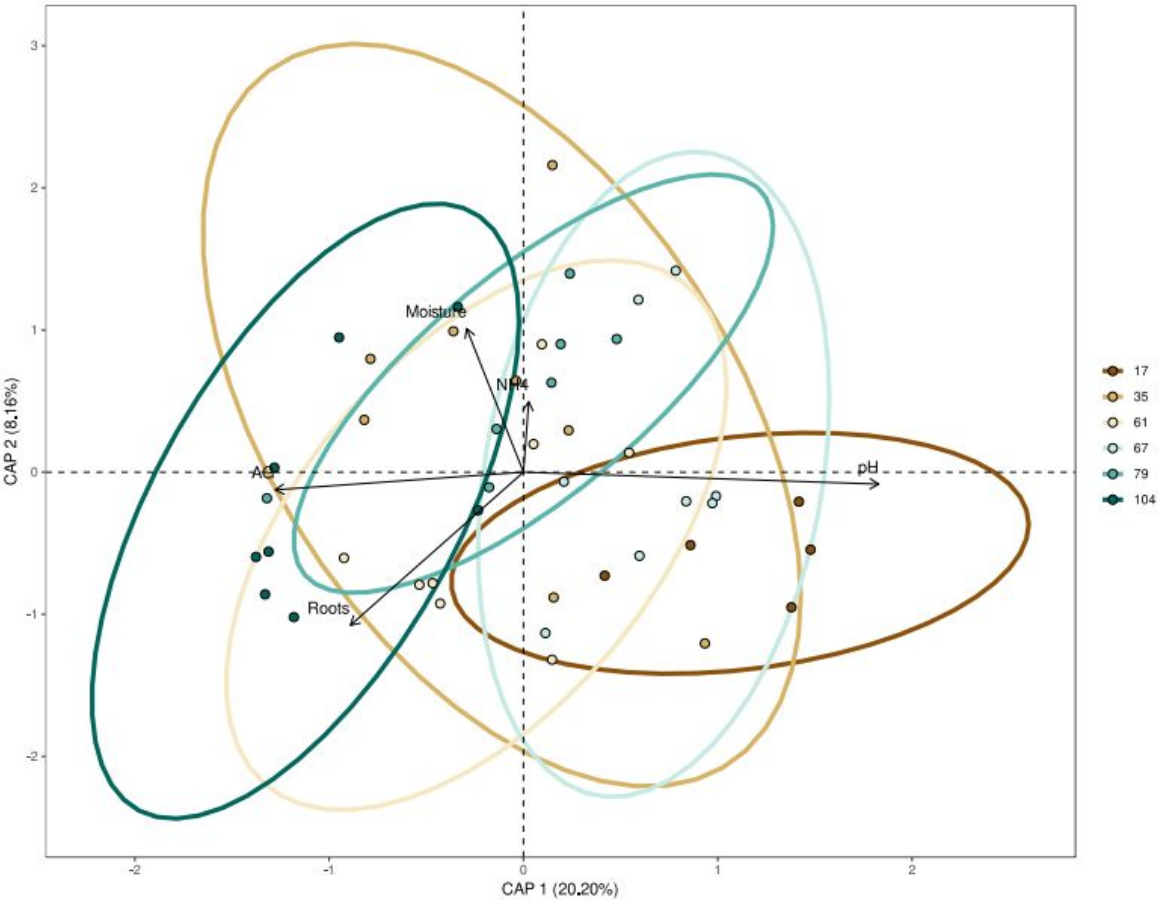
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1340 **Figure 9**



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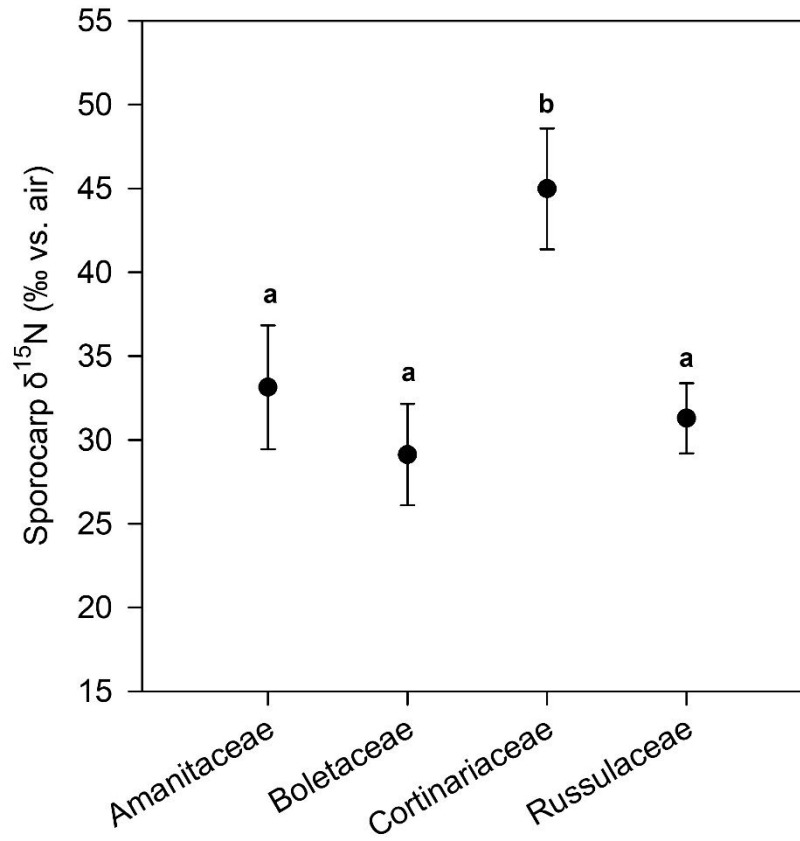
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1349 **Figure 10**

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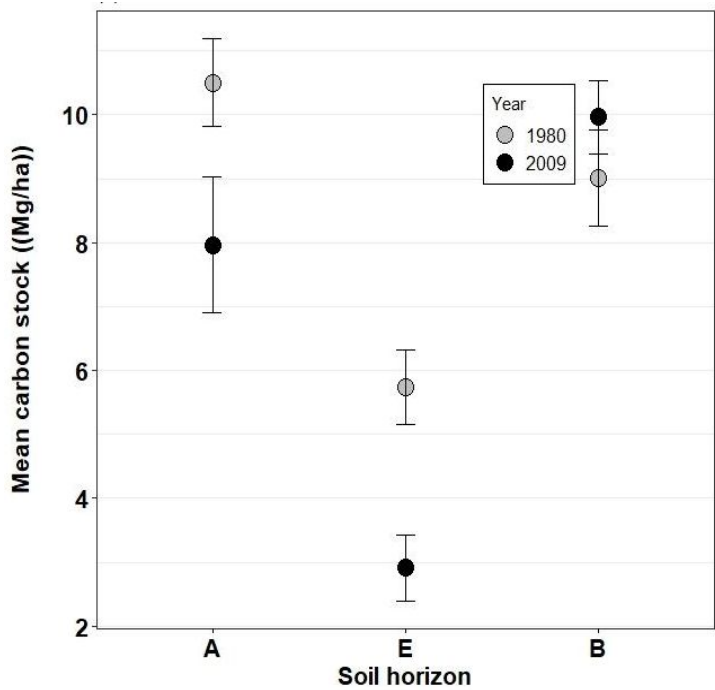
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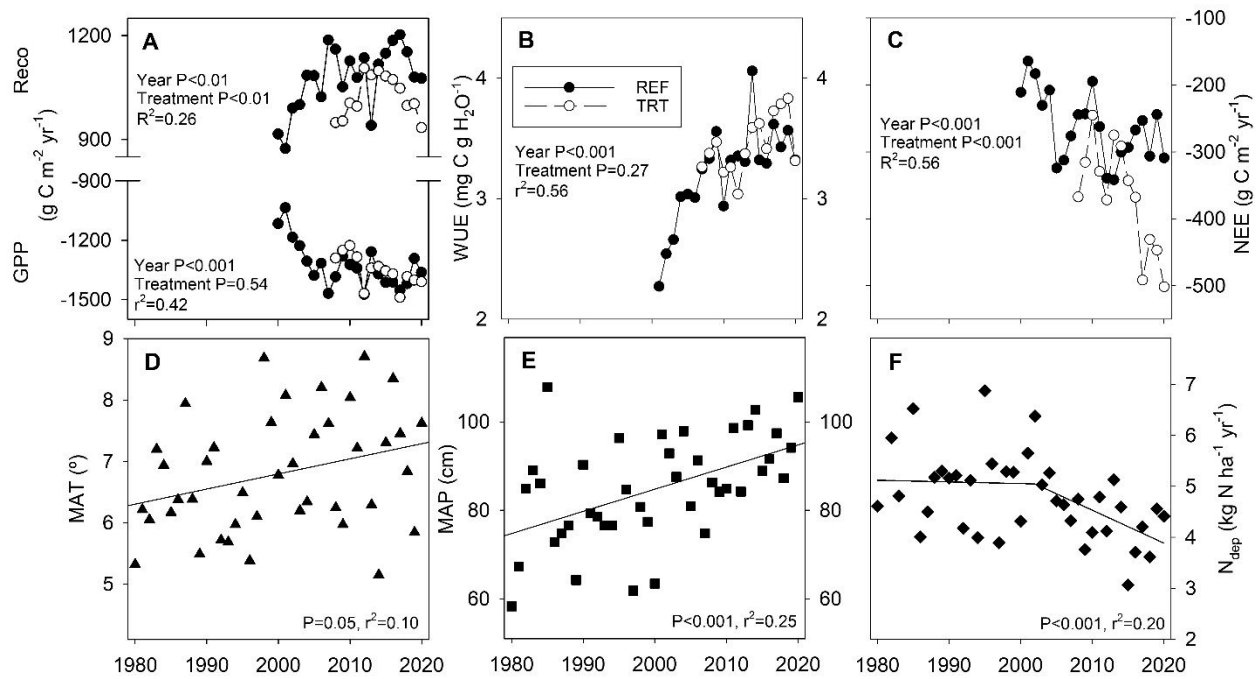
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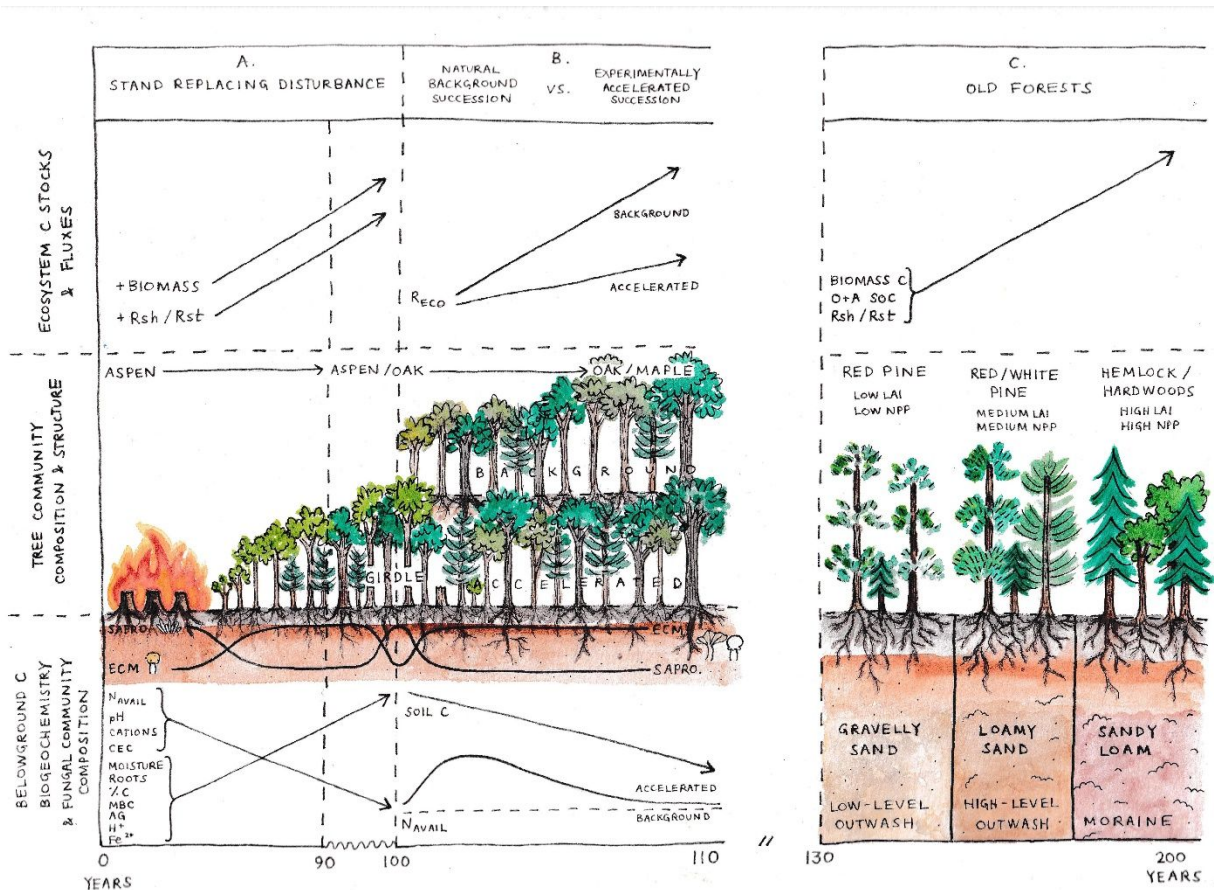
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Figure 11



1369 **Figure 12**

1380 **Figure 13**



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Appendix S1

Lucas Nave, Chris Gough, Cameron Clay, Fernanda Santos, Jeff Atkins, Sonja Benjamins-Carey, Gil Bohrer, Buck Castillo, Robert Fahey, Brady Hardiman, Kathryn Hofmeister, Valeriy Ivanov, Jennifer Kalejs, Ashley Matheny, Angela Menna, Knute Nadelhoffer, Brooke Propson, Adam Schubel, Jason Tallant

Carbon cycling across ecosystem succession in a north temperate forest: controls and management implications. *Ecological Applications*.

1. Detailed Methods

Meta-analysis of organic horizon C stocks across the Burn Plots chronosequence- In order to test for disturbance effects on O horizon C stocks over successional timescales, we synthesized data from 5 known soil sampling campaigns on the Burn Plots, including Schaetzl (1994; sampled 1992), Gough et al. (2007; sampled 2004), and Nave et al. (2019; sampled 2014), and previously unpublished data from collections in 1993 and 2012. The 1993 dataset was collected and documented in a class project (Lehman et al. 1993); the 2012 dataset was collected and documented by a longtime close collaborator to many of the coauthors of this synthesis (Le Moine 2012). Across the five sampling campaigns, frames were used to collect fixed-area O horizon samples, down to the surface of the underlying mineral soil. Fixed-area samples were dried, weighed, ground, and analyzed either for %C (in the case of the three published papers), or, in the case of the unpublished datasets, by loss on ignition, which we multiplied by an assumed C fraction of 0.5 to calculate O horizon C stocks in Mg ha^{-1} as the product of C concentration and dry mass per area. Assuming differences in individual judgment, sampling techniques, and other factors, we chose to use unweighted effect-size meta-analysis to analyze

the data, following the methods of Nave et al. (2021). We computed the response ratio for each experimental treatment stand as its mean O horizon C stock divided by the mean O horizon C stock of the oldest stand in the chronosequence that was sampled at that time. This approach was necessary because individual sampling campaigns sampled different portions of the chronosequence at different times in its history. Collectively, this dataset (Appendix S2) compares chronosequence stands that were 6-56, 18-57, 12-81, 14-76, or 16-78 years old to stands that were 68, 82, 90, 101, or 103 years old when sampled.

Soil respiration scaling- Methods for scaling instantaneous point measurements of R_s to annual and growing season C fluxes in the chronosequence and old reference forest stands are documented in Clay et al. (2022). For scaling point measurements to cumulative fluxes in the tower footprints, where R_s was measured in the 1.1 ha intensive plot surrounding each tower, we first developed temperature-response functions for each footprint (UMB and UMd), utilizing the full span of available R_s measurements (1998-2020 for UMB, 2008-2017 for UMd). These functions were exponential models relating instantaneous R_s to soil temperature, of the form $R_s = a * e^{(b * T_s)}$, where a and b were footprint-specific parameters and T_s was the simultaneously measured soil temperature at 7.5 cm depth. Next, we used continuous hourly (1998-2006) or half-hourly (2007-present) sensor-based T_s measurements (7.5 cm depth), accessed via AmeriFlux (<https://ameriflux.lbl.gov/data/download-data/>) to compute cumulative annual R_s efflux, scaled to $Mg\ C\ ha^{-1}$ based on collar size and within-plot replication. We only performed these computations for years in which >95% of the T_s data were either measured, or were gap-filled with high confidence (Pastorello et al. 2020). This eliminated 2015-16 from UMB and 2015 from UMd.

Longitudinal sampling of soil C stocks- To assess changes in mineral soil C stocks over recent decades, soil was collected in 1980 and 2009 in an unmanipulated control area adjacent to the Burn Plots. This area consists of 1911-origin forestland that established after the cutting and burning disturbance that was widespread across the UMBS landscape and the wider region. In 1980, a sampling plot with dimensions of 100m x 120m was established, and soils were sampled by genetic horizon (A, E, and Bs1) at sampling points equally spaced along 3 transects. Each soil horizon composite consisted of five individual soil samples collected using a soil corer (0-30cm depth) and at 1-meter radius around each sampling point. In 2009, the same sampling design was implemented approximately 30 m from the perimeter of the 1980 sampling plot. At both campaigns, soils were air-dried and sieved (< 2 mm mesh size), oven dried at 60°C until they reached constant mass, and fine earth bulk densities were measured and used to calculate soil C stocks in Mg C ha⁻¹ for each genetic horizon.

Soil biogeochemistry and microbial communities- To characterize long-term successional patterns in soil physicochemical, biogeochemical, microbial properties, we report data from a systematic 2015 sampling campaign on the Burn Plots (Nave et al. 2024). In each Burn Plot (i.e., experimental stand), 5 A horizon samples were collected and composited at each of 3 locations, using a 5.2 cm diameter core, and stored at 4° until processing (within 3 wk). The resulting 3 composite samples per stand were sieved (2mm mesh) to remove rocks and roots from the fine earth fraction. Roots were oven dried at 60° for 24 hours and multiplied by core area to determine root biomass density. Immediately after sieving, the fine earth was subsampled for DNA sequencing, extracellular enzyme activity, microbial biomass C and soil N availability, solid-phase C and N concentrations, pH, and gravimetric water content.

68 Subsamples for DNA sequencing (2 g) were flash-frozen in liquid nitrogen and stored at -80 C
69 until extraction. Genomic DNA was extracted using a Powersoil DNA Elution Kit (MOBIO,
70 Carlsbad, CA USA), extraction quantity determined using a Qubit Fluorometer (Invitrogen, San
71 Jose, CA, USA) and extracts stored at -80° until amplification. The ITS2 region was amplified
72 using forward primer 5.8S-FUN (5'-AGWGATCCRTTGYYRAAAGTTCCCTGACTGACT)
73 and reverse primer ITS4-FUN (GCAWAWCAAWAAGCGGAGGACCCTGACTGACT). Each
74 subsample had paired forward and reverse barcodes appended to the primer ends for
75 multiplexing. PCR reactions were carried out using Phusion High-Fidelity DNA Polymerase
76 (New England BioLabs, Ipswich, MA) under the following conditions: 5.9ul PCR grade H₂O,
77 4ul 5X Phusion HF Buffer, 0.4ul 10mM dNTP mix, 0.75ul forward primer (10mM), 0.75ul
78 reverse primer (10mM), 8ul DNA template, 0.2ul Phusion Taq. PCR cycle parameters were as
79 follows: 3 min of initial denaturation at 94°, 28 cycles of 30 sec denaturation at 94°, 45 sec
80 annealing at 55°, and 90 sec at 72° with a final extension at 72° for 10 min. PCR products were
81 submitted to the Microbial Systems Molecular Biology Laboratory at the University of Michigan
82 for sequencing and demultiplexing. Sequences were processed using QIIME2 package version
83 qiime2-2019.4. Prior to downstream analysis, forward and reverse reads were analyzed for QC
84 scores in QIIME 2 View. All reverse reads were excluded due to poor quality and only forward
85 reads were used in downstream analyses. Sequences were trimmed for poor quality end reads
86 using denoise-single trimming the first 12 bp and truncating at 245 bp. Sequences were clustered
87 into OTUs at 97% similarity and filtered for chimeras using a vsearch cluster-features-de-novo
88 pipeline. OTUs were then classified using a classifier created from the UNITE database using the
89 classify-sklearn command. A phylogenetic tree was generated with QIIME phylogeny fasttree.
90 The dataset was then filtered for sequences that appeared less than 5 times and fewer than 2

91 samples with filter-features and rarefied to 6845, representing the minimum sequences total for
92 an individual sample. The most frequent OTUs are presented in Table S1.

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For Review Only

99 **Table S1.** Fungal OTUs individually accounting for >1% relative abundance of total sequences. Trophic status was assigned to OTUs
100 classified (>97%) at the genus level.

Trophic status	Family	Genus	Species	% relative abundance	Total sequence #
Ectomycorrhizal	Gloniaceae	Cenococcum	<i>Cenococcum geophilum</i>	5.7	35496
Ectomycorrhizal	Cortinariaceae	Cortinarius	<i>Cortinarius alpinus</i>	5.4	33752
Ericoid mycorrhizal	Helotiaceae	Meliniomyces	<i>Meliniomyces bicolor</i>	3.3	20523
Ectomycorrhizal	Russulaceae	Lactifluus	<i>Lactifluus allardii</i>	3.1	19456
Ectomycorrhizal	Amanitaceae	Amanita	<i>Amanita brunnescens</i>	2.9	18217
Ectomycorrhizal	Cortinariaceae	Cortinarius	<i>Cortinarius caperatus</i>	2.8	17664
Unidentified	Russulaceae	unidentified	<i>unidentified</i>	2.7	17104
Ectomycorrhizal	Hydnangiaceae	Laccaria	<i>unidentified</i>	2.3	14509
Ectomycorrhizal	Atheliaceae	Piloderma	<i>Piloderma bicolor</i>	2.0	12364
Ectomycorrhizal	Cortinariaceae	Cortinarius	<i>unidentified</i>	1.6	9770
Unidentified	unidentified	unidentified	<i>unidentified</i>	1.4	8716
Endophytic	Umbelopsidaceae	Umbelopsis	<i>Umbelopsis dimorpha</i>	1.3	8379
unidentified	unidentified	unidentified	<i>unidentified</i>	1.3	8055
Ectomycorrhizal	Russulaceae	Russula	<i>unidentified</i>	1.3	7883
Ectomycorrhizal	Cortinariaceae	Cortinarius	<i>Cortinarius cruentiphylus</i>	1.2	7590
unidentified	Russulaceae	unidentified	<i>unidentified</i>	1.2	7487
Saprobic	Omphalotaceae	Rhodocollybia	<i>Rhodocollybia butyracea</i>	1.2	7273
Ectomycorrhizal	Cortinariaceae	Cortinarius	<i>Cortinarius leiocastaneus</i>	1.1	6763
Saprobic	Clavariaceae	Clavulinopsis	<i>Clavulinopsis laeticolor</i>	1.1	6749
unidentified	Herpotrichiellaceae	unidentified	<i>unidentified</i>	1.1	6580
Ectomycorrhizal	Cortinariaceae	Cortinarius	<i>Cortinarius alboviolaceus</i>	1.1	6537
Ectomycorrhizal	Cantharellaceae	Craterellus	<i>Craterellus tubaeformis</i>	1.0	6378

102 Soil extracellular enzyme activity, microbial biomass C, and available N determinations
103 followed standard methods. For extracellular enzyme activity (Sinsabaugh and Linkins 1990),
104 three 1 g subsamples from each fine earth sample were extracted in 125 mL of 50mM sodium
105 acetate buffer (pH 5) and α -glucosidase activity was measured after a 4-hour incubation with 4-
106 nitrophenyl α -D-glucopyranoside on a Biotek ELX800 plate reader (Biotek, Winooski, VT,
107 USA). For microbial biomass C (Wu et al. 1990), two 4g subsamples from each fine earth
108 sample were weighed into 15ml polypropylene vials. One subsample was extracted while the
109 other was placed in a sterilized 125 ml Erlenmeyer flask with ~20 boiling chips and 75 ml of
110 ethanol-free chloroform in a desiccator for 48 hours. Both subsamples were extracted for 1 hour
111 with 0.1 M K₂SO₄ (5:1 solution: soil ratio) on a shaker table (180 strokes per minute). Extracts
112 were filtered, mixed with 15 ml of sodium hexametaphosphate and 2% potassium persulfate, and
113 analyzed for total organic C (TOC) on a Seal AA3 autoanalyzer. Microbial biomass C was
114 calculated as the difference in TOC between non-fumigated and fumigated soils. Soil available N
115 was measured on a 3 g subsample of fine earth as [NH₄-N], following extraction with 30 mL of
116 2 M KCl and quantitation on a Seal AA3 autoanalyzer according to U.S. EPA method 350.1.
117 Concentrations of TOC and NH₄-N were scaled to an oven-dry soil basis (mass per mass) using
118 the relevant volumes of extractants and reactants, measured masses of extracted subsamples, and
119 gravimetric moisture content. Alpha glucosidase activity rates were scaled to the quantity of
120 substrate consumed per unit time per mass of soil.

121 Fine earth subsamples prepared for pH (~5 g, 1:1 soil:water slurry) were measured using a model
122 8000 pH meter (VWR Scientific, Radnor, PA, USA), and gravimetric moisture content and %C
123 after lyophilization (5 g), with elemental analysis on a Costech ECS 4010 elemental analyzer in
124 the UMBS Analytical Laboratory. The UMBS Analytical Laboratory also housed a Finnigan

Delta V isotope ratio mass spectrometer, which was used to analyze the ^{15}N signatures of ectomycorrhizal (EcM) fungal sporocarps collected in July 2022 from paired plots (n=6 plots) in the reference and treatment flux tower footprints that received ^{15}N tracer additions in 2010 (Nave et al. 2013). We analyzed these fungal samples in order to assess long-term N foraging of several EcM fungal families well represented in Table S1, using ^{15}N -enrichment as an indicator.

Climatology and atmospheric deposition- Daily climatology measurements and weekly atmospheric deposition collections have occurred since the late 1970s on the main “campus” of UMBS, situated on the south shore of 1,700 ha Douglas Lake. The weather station is not inspected by the National Weather Service cooperative network but meets the design standards of its installations, with daily maximum and minimum air temperatures at 2 m height recorded on maximum and minimum Liquid In Glass thermometers, inside a Cotton Region Shelter. Daily water-equivalent precipitation is measured at the UMBS National Atmospheric Deposition Program (NADP) site (site MI09, 45.5608, -84.6783) via a Belfort Universal Precipitation Gage 5-780 (July 1, 1979 to January 7, 2020) or an ETI NOAA IV Total Precipitation Gauge (January 8, 2020 to present). In this paper we present mean annual temperatures as the averages of daily maximum and minimum values, and mean annual precipitation as the sum of daily water-equivalent values. Wet atmospheric deposition is collected weekly in an open field with an Aerochem Metrics Precipitation Collector according to standard NADP protocols (National Trends Network 2019). Weekly samples were mailed to the Central Analytical Laboratory at the University of Illinois (July 1979 to February 2018) or the University of Wisconsin State Lab of Hygiene (March 2018 to present) for analysis. In this paper we accessed NADP National Trends Network data to present annual total inorganic N deposition in $\text{kg N ha}^{-1} \text{ yr}^{-1}$.

Data analysis- The results presented in this study are based on community-standard approaches and tests. In most analyses, we used chronosequence stands or flux tower plots as the experimental units, e.g., to test for changes in C pools over time, or to quantify relationships between C fluxes and drivers. For some analyses aimed at uncovering more mechanistic relationships, spatial variation within chronosequence stands facilitated the use of individual plots as experimental units. To analyze tower-based fluxes, we treated the two footprints as experimental units. Table S2 provides, for each relationship tested using inferential statistics, information about which units of observation were treated as the experimental units, the response and predictor variables used in the statistical test, and information about the statistical test. This tabular detail is offered as supplementary to the data analysis section (2.4) and results narrative (section 3) of the main paper. After table S2, further detail is provided in section 2 of this Appendix in the form of statistical test outputs.

Table S2. Summary of relationships tested using inferential statistics.

Relationship	Experimental Units	Response Variable	Predictor Variable(s)	Test	Additional Notes
Ecosystem C vs. stand age	Individual plots in each chronosequence & old reference stand, binned into 40-yr age classes	Individual pool & total ecosystem C stocks	Age class	ANOVA	ANOVA of C stock in each pool, by plot (n=30 total)
Canopy structure vs. C uptake	Individual plots in each chronosequence & old reference stand	Wood NPP	LAI, canopy height, rugosity	multiple linear regression	One regression each for young (22 - 48 yr), middle (66-84), & old (>100) stands. Predictor variables standardized (subtract from mean, divide by stdev) prior to analysis. Predictive capacity of individual variables assessed by partial r^2 values.
Leaf production over time, by taxon	Individual plots in reference and treatment tower footprints	Annual leaf litterfall flux	Year, footprint, taxon	simple linear regression	Separate regressions for each species, genus, or functional group and footprint
Soil respiration vs. stand age	Individual chronosequence & old reference stands	Rst, Rsa, Rsh, Rsh/Rst	Stand age	simple linear regression	Separate regressions for annual and growing season means for each response variable
Fine root production, stand age, and soil respiration	Individual chronosequence & old reference stands	Growing season root production; annual Rst	Stand age; growing season root production	simple linear regression	Two separate regressions, all values averaged across all available years of observation

Table S2 continued.

Relationship	Experimental Units	Response Variable	Predictor Variable(s)	Test	Additional Notes
O horizon C stocks vs. stand age	Individual chronosequence stands	O horizon C stock	Treatment, Stand age	unweighted meta-analysis	Categorical meta-analysis to test for disturbance effect; tested for significant temporal pattern with continuous meta-analysis
Soil properties vs. stand age	Individual chronosequence stands	%C, microbial biomass C, and pH	Stand age	simple linear regression	A horizons, data from 2x cut+burn stands only, sampled in 2015
Soil enzymes vs. properties	Individual chronosequence stands	α -glucosidase activity rate	%C, pH	simple linear regression	A horizons, data from 2x cut+burn stands only, sampled in 2015
Fungal community composition vs. stand age	Soil samples from individual chronosequence stands	Bray-Curtis community dissimilarity	principal coordinate axes	Principal coordinates analysis	Mantel test of significant differences in community dissimilarity across stand ages
Fungal community composition vs. stand age and soil properties	Soil samples from individual chronosequence stands	Distance-based dissimilarity values	pH, %C, %N α -glucosidase, NH ₄ , soil moisture, fine root biomass	dbRDA, forward stepwise regression	Forward stepwise regression to identify significant predictors of variation in community dissimilarity

Table S2 continued.

Relationship	Experimental Units	Response Variable	Predictor Variable(s)	Test	Additional Notes
Fungal N foraging and competition	Individual plots in reference and treatment tower footprints	Sporocarp 15N enrichment	EcM family	ANOVA	ANOVA of 15N values of bulk sporocarps in each family from each plot
Soil C stock change in recent decades	Individual soil profiles from common control area	A, E, B horizon C stocks	Year, Horizon	2-way ANOVA	Samples collected in 1980 vs. 2009, within 30 m distance of each other, processed and analyzed using common methodology.
C uptake and losses	Reference and treatment tower footprints	Annual GPP, Reco, NEE	Year, treatment	multiple linear regression	One regression for each response variable, with year (continuous) and treatment (dummy) as predictors
Canopy water use efficiency	Reference and treatment tower footprints	Growing season WUE	Year, treatment	multiple linear regression	Year (continuous) and treatment (dummy) as predictors
Climate change	Weather station	MAT, MAP	Year	simple linear regression	Simple linear regressions to test for significant slope (change) in climate variables over time
N deposition over time	NADP station	Total inorganic N	Year	piecewise regression	Regression to test for significant change in N deposition over time

2. Statistical output

This section provides the output of the statistical tests underpinning each of the figures in the main manuscript, except for Figures 8 and 9. Data and code (QIIME) for the statistical analyses underlying those figures are in Nave et al (2024).

Figure 2a: Ecosystem C stocks vs. stand age

One Way Analysis of Variance

Wednesday, February 06, 2019, 11:52:08 AM

Data source: Data 3 in C pools stats

Dependent Variable: Total ecosystem (Mg C ha⁻¹)

Normality Test (Shapiro-Wilk) Passed (P = 0.751)

Equal Variance Test: Passed (P = 0.400)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	106.927	19.231	6.799
40-80	9	0	134.753	10.911	3.637
80-120	3	0	163.911	7.137	4.121
120-160	6	0	199.419	24.362	9.946
>160	3	0	318.182	14.476	8.358

Source of Variation	DF	SS	MS	F	P
Between Groups	4	112530.034	28132.509	96.045	<0.001
Residual	24	7029.862	292.911		
Total	28	119559.896			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = <0.001).

Power of performed test with alpha = 0.050: 1.000

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: **Age class**

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
>160 vs. <40	211.254	23.914	<0.001	Yes
>160 vs. 40-80	183.428	23.549	<0.001	Yes
>160 vs. 80-120	154.271	28.841	<0.001	Yes
>160 vs. 120-160	118.762	24.977	<0.001	Yes
120-160 vs. <40	92.492	19.077	<0.001	Yes
120-160 vs. 40-80	64.666	18.617	<0.001	Yes
120-160 vs. 80-120	35.509	24.977	0.007	Yes
80-120 vs. <40	56.983	23.914	<0.001	Yes
80-120 vs. 40-80	29.158	23.549	0.017	Yes
40-80 vs. <40	27.826	17.164	0.003	Yes

Wednesday, February 06, 2019, 11:46:14 AM

One Way Analysis of Variance

Data source: Data 2 in C pools stats

Dependent Variable: Live biomass total (Mg C ha-1)

Normality Test (Shapiro-Wilk) Passed (P = 0.426)

Equal Variance Test: Passed (P = 0.322)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	31.979	8.593	3.038
40-80	9	0	57.849	8.559	2.853
80-120	3	0	81.212	3.850	2.223
120-160	6	0	117.000	14.595	5.958
>160	3	0	224.343	18.597	10.737

Source of Variation	DF	SS	MS	F	P
Between Groups	4	93392.159	23348.040	193.936	<0.001
Residual	24	2889.372	120.390		
Total	28	96281.531			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = <0.001).

Power of performed test with alpha = 0.050: 1.000

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: Age class

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
>160 vs. <40	192.363	15.331	<0.001	Yes
>160 vs. 40-80	166.494	15.097	<0.001	Yes
>160 vs. 80-120	143.131	18.490	<0.001	Yes
>160 vs. 120-160	107.343	16.013	<0.001	Yes
120-160 vs. <40	85.020	12.230	<0.001	Yes
120-160 vs. 40-80	59.151	11.935	<0.001	Yes
120-160 vs. 80-120	35.788	16.013	<0.001	Yes
80-120 vs. <40	49.232	15.331	<0.001	Yes
80-120 vs. 40-80	23.363	15.097	0.004	Yes
40-80 vs. <40	25.869	11.004	<0.001	Yes

One Way Analysis of Variance

Wednesday, February 06, 2019, 11:47:26 AM

Data source: Data 2 in C pools stats

Dependent Variable: sqrt_STD_DEAD

Normality Test (Shapiro-Wilk) Passed (P = 0.634)**Equal Variance Test:** Passed (P = 0.224)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	1.293	0.308	0.109
40-80	9	0	1.882	0.361	0.120
80-120	3	0	2.474	0.583	0.337
120-160	6	0	1.358	0.506	0.207
>160	3	0	1.090	0.0684	0.0395

Source of Variation	DF	SS	MS	F	P
Between Groups	4	4.890	1.222	7.987	<0.001
Residual	24	3.673	0.153		
Total	28	8.563			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = <0.001).

Power of performed test with alpha = 0.050: 0.986

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: **Age class**

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
80-120 vs. >160	1.384	0.659	<0.001	Yes
80-120 vs. <40	1.181	0.547	<0.001	Yes
80-120 vs. 120-160	1.116	0.571	<0.001	Yes
80-120 vs. 40-80	0.592	0.538	0.033	Yes
40-80 vs. >160	0.792	0.538	0.006	Yes
40-80 vs. <40	0.589	0.392	0.005	Yes
40-80 vs. 120-160	0.524	0.426	0.018	Yes
120-160 vs. >160	0.268	0.571	0.342	No
120-160 vs. <40	0.0647	0.436	0.762	Do Not Test
<40 vs. >160	0.204	0.547	0.450	Do Not Test

One Way Analysis of Variance

Wednesday, February 06, 2019, 11:53:36 AM

Data source: Data 2 in C pools stats

Dependent Variable: DWD Mg C ha-1

Normality Test (Shapiro-Wilk) Passed (P = 0.059)

Equal Variance Test: Passed (P = 0.059)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	1.700	1.335	0.472
40-80	9	0	1.369	0.765	0.255
80-120	3	0	1.839	0.530	0.306
120-160	6	0	0.872	0.928	0.379
>160	3	0	6.332	3.838	2.216

Source of Variation	DF	SS	MS	F	P
Between Groups	4	68.123	17.031	7.940	<0.001
Residual	24	51.480	2.145		
Total	28	119.603			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = <0.001).

Power of performed test with alpha = 0.050: 0.985

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: Age class

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
>160 vs. 120-160	5.460	2.137	<0.001	Yes
>160 vs. 40-80	4.963	2.015	<0.001	Yes
>160 vs. <40	4.631	2.046	<0.001	Yes
>160 vs. 80-120	4.493	2.468	<0.001	Yes
80-120 vs. 120-160	0.967	2.137	0.360	No
80-120 vs. 40-80	0.470	2.015	0.635	Do Not Test
80-120 vs. <40	0.138	2.046	0.890	Do Not Test
<40 vs. 120-160	0.828	1.632	0.305	Do Not Test
<40 vs. 40-80	0.332	1.469	0.645	Do Not Test
40-80 vs. 120-160	0.497	1.593	0.526	Do Not Test

One Way Analysis of Variance

Wednesday, February 06, 2019, 10:34:48 AM

Data source: Data 2 in C pools stats

Dependent Variable: O+A horiz Mg C ha-1

Normality Test (Shapiro-Wilk) Passed (P = 0.058)**Equal Variance Test:** Passed (P = 0.550)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	17.407	5.929	2.096
40-80	9	0	18.066	5.410	1.803
80-120	3	0	20.781	3.220	1.859
120-160	6	0	28.416	13.118	5.355
>160	3	0	34.103	6.082	3.512

Source of Variation	DF	SS	MS	F	P
Between Groups	4	998.886	249.722	4.175	0.010
Residual	24	1435.368	59.807		
Total	28	2434.254			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = 0.010).

Power of performed test with alpha = 0.050: 0.739

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: **Age class**

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
>160 vs. <40	16.696	10.806	0.004	Yes
>160 vs. 40-80	16.037	10.641	0.005	Yes
>160 vs. 80-120	13.322	13.032	0.045	Yes
>160 vs. 120-160	5.687	11.286	0.309	No
120-160 vs. <40	11.009	8.620	0.014	Yes
120-160 vs. 40-80	10.350	8.412	0.018	Yes
120-160 vs. 80-120	7.636	11.286	0.175	No
80-120 vs. <40	3.373	10.806	0.526	No
80-120 vs. 40-80	2.714	10.641	0.603	Do Not Test
40-80 vs. <40	0.659	7.756	0.862	Do Not Test

One Way Analysis of Variance

Wednesday, February 06, 2019, 10:37:23 AM

Data source: Data 2 in C pools stats

Dependent Variable: Fine root Mg C ha-1

Normality Test (Shapiro-Wilk) Passed (P = 0.648)

Equal Variance Test: Passed (P = 0.772)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	6.539	1.956	0.692
40-80	9	0	7.778	1.533	0.511
80-120	3	0	7.944	0.579	0.334
120-160	6	0	5.917	2.480	1.012
>160	3	0	4.154	1.377	0.795

Source of Variation	DF	SS	MS	F	P
Between Groups	4	38.437	9.609	2.854	0.046
Residual	24	80.799	3.367		
Total	28	119.236			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = 0.046).

Power of performed test with alpha = 0.050: 0.473

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: Age class

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
80-120 vs. >160	3.791	3.092	0.018	Yes
80-120 vs. 120-160	2.027	2.678	0.131	No
80-120 vs. <40	1.405	2.564	0.269	Do Not Test
80-120 vs. 40-80	0.166	2.525	0.893	Do Not Test
40-80 vs. >160	3.624	2.525	0.007	Yes
40-80 vs. 120-160	1.861	1.996	0.066	Do Not Test
40-80 vs. <40	1.238	1.840	0.178	Do Not Test
<40 vs. >160	2.386	2.564	0.067	No
<40 vs. 120-160	0.623	2.045	0.536	Do Not Test
120-160 vs. >160	1.763	2.678	0.187	Do Not Test

One Way Analysis of Variance

Wednesday, February 06, 2019, 10:36:57 AM

Data source: Data 2 in C pools stats

Dependent Variable: Minsoil 0-100 Mg C ha-1

Normality Test (Shapiro-Wilk) Passed (P = 0.952)**Equal Variance Test:** Passed (P = 0.362)

Group Name	N	Missing	Mean	Std Dev	SEM
<40	8	0	47.545	10.585	3.742
40-80	9	0	46.034	3.170	1.057
80-120	3	0	45.788	6.348	3.665
120-160	6	0	45.157	11.554	4.717
>160	3	0	48.059	4.609	2.661

Source of Variation	DF	SS	MS	F	P
Between Groups	4	30.262	7.566	0.110	0.978
Residual	24	1655.114	68.963		
Total	28	1685.376			

Figure 3: Canopy complexity and NPP_w

Multiple Linear Regression

Tuesday, November 21, 2023 11:39:46 AM

Data source: Data 1 in Notebook3

Young NPPkgC/ha = -386.904 + (212.141 * LAI 4 Ring) + (165.573 * mean.max.ht) - (122.763 * rugosity)

N = 11

R = 0.680 Rsqr = 0.462 Adj Rsqr = 0.232

Standard Error of Estimate = 344.238

	Coefficient	Std. Error	t	P	VIF
Constant	-386.904	786.526	-0.492	0.638	
LAI 4 Ring	212.141	325.718	0.651	0.536	1.089
mean.max.ht	165.573	95.641	1.731	0.127	4.918
rugosity	-122.763	125.903	-0.975	0.362	4.771

Warning: Multicollinearity is present among the independent variables. The variables with the largest values of VIF are causing the problem. Consider getting more data or eliminating one or more variables from the equation. The likely candidates for elimination are: mean.max.ht , rugosity

Analysis of Variance:

	DF	SS	MS	F	P
Regression	3	712988.378	237662.793	2.006	0.202
Residual	7	829499.816	118499.974		
Total	10	1542488.194	154248.819		

Multiple Linear Regression

Tuesday, November 21, 2023 11:49:45 AM

Data source: Data 1 in complexity.JNB

Mid NPP = 929.574 - (347.274 * LAI) + (80.977 * Max) + (9.149 * Rug)

N = 6

R = 0.989 Rsqr = 0.978 Adj Rsqr = 0.945

Standard Error of Estimate = 18.296

	Coefficient	Std. Error	t	P	VIF
Constant	929.574	141.049	6.590	0.022	
LAI	-347.274	57.516	-6.038	0.026	2.302
Max	80.977	11.407	7.099	0.019	1.797
Rug	9.149	4.701	1.946	0.191	1.435

Analysis of Variance:

	DF	SS	MS	F	P
Regression	3	30031.531	10010.510	29.904	0.033
Residual	2	669.513	334.757		
Total	5	30701.045	6140.209		

Multiple Linear Regression

Tuesday, November 21, 2023 11:50:11 AM

Data source: Data 1 in complexity.JNB

Old NPP = -299.893 + (1108.522 * LAI) + (9.847 * Max) - (38.737 * Rug)

N = 12

R = 0.823 Rsqr = 0.678 Adj Rsqr = 0.557

Standard Error of Estimate = 647.868

	Coefficient	Std. Error	t	P	VIF
Constant	-299.893	1933.172	-0.155	0.881	
LAI	1108.522	281.843	3.933	0.004	1.009
Max	9.847	128.494	0.0766	0.941	3.345
Rug	-38.737	47.973	-0.807	0.443	3.354

Analysis of Variance:

	DF	SS	MS	F	P
Regression	3	7072270.398	2357423.466	5.616	0.023
Residual	8	3357863.244	419732.905		
Total	11	10430133.641	948193.967		

Figure 4: Leaf litterfall compositional change

Linear Regression

Wednesday, May 3, 2023 2:59:38 PM

Data source: summaries in Myco_succession_litterfall.JNB

REF POP+BEPA = 21211.587 - (10.256 * Year)

N = 23

R = 0.760 Rsqr = 0.578 Adj Rsqr = 0.558

Standard Error of Estimate = 60.794

	Coefficient	Std. Error	t	P
Constant	21211.587	3837.366	5.528	<0.001
Year	-10.256	1.911	-5.367	<0.001

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	106452.737	106452.737	28.803	<0.001
Residual	21	77612.942	3695.854		
Total	22	184065.680	8366.622		

Normality Test (Shapiro-Wilk) Passed (P = 0.051)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.061)

Linear Regression

Wednesday, May 3, 2023 1:50:13 PM

Data source: summaries in Myco_succession_litterfall.JNB

REF QURU = -9152.636 + (4.653 * Year)

N = 23

R = 0.857 Rsqr = 0.735 Adj Rsqr = 0.722

Standard Error of Estimate = 19.393

	Coefficient	Std. Error	t	P
Constant	-9152.636	1224.129	-7.477	<0.001
Year	4.653	0.610	7.633	<0.001

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	21911.740	21911.740	58.261	<0.001
Residual	21	7898.086	376.099		
Total	22	29809.826	1354.992		

Normality Test (Shapiro-Wilk) Passed (P = 0.127)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.630)

617

618 **Linear Regression**

Wednesday, May 3, 2023 2:04:42 PM

619

620 **Data source:** summaries in Myco_succession_litterfall.JNB

621

622 $\text{TRT QURU} = -29196.253 + (14.699 * \text{Year})$

623

624 $N = 14$

625

626 $R = 0.802$ $\text{Rsqr} = 0.643$ $\text{Adj Rsqr} = 0.613$

627

628 Standard Error of Estimate = 47.735

629

	Coefficient	Std. Error	t	P
Constant	-29196.253	6369.192	-4.584	<0.001
Year	14.699	3.165	4.644	<0.001

633

634 Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	49153.125	49153.125	21.571	<0.001
Residual	12	27343.732	2278.644		
Total	13	76496.857	5884.374		

639

640 Normality Test (Shapiro-Wilk) Passed (P = 0.894)

641

642 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.904)

643

644 **Linear Regression**

Wednesday, May 3, 2023 2:04:55 PM

645

646 **Data source:** summaries in Myco_succession_litterfall.JNB

647

648 $\text{TRT Acer} = -10353.956 + (5.332 * \text{Year})$

649

650 $N = 14$

651

652 $R = 0.571$ $\text{Rsqr} = 0.326$ $\text{Adj Rsqr} = 0.269$

653

654 Standard Error of Estimate = 33.409

655

	Coefficient	Std. Error	t	P
Constant	-10353.956	4457.676	-2.323	0.039
Year	5.332	2.215	2.407	0.033

659

660 Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	6467.556	6467.556	5.794	0.033
Residual	12	13393.873	1116.156		
Total	13	19861.429	1527.802		

665

666 Normality Test (Shapiro-Wilk) Passed (P = 0.793)

667

668 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.715)

Figure 5: Soil respiration vs stand age

Linear Regression

Wednesday, May 22, 2024 11:27:25 AM

Data source: Data 1 in synthesis figs.JNB

$$RshA/RstA = 0.473 + (0.00118 * Stand\ Age)$$

N = 11

R = 0.605 Rsqr = 0.366 Adj Rsqr = 0.295

Standard Error of Estimate = 0.084

	Coefficient	Std. Error	t	P
Constant	0.473	0.0501	9.451	<0.001
Stand Age	0.00118	0.000517	2.279	0.049

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	0.0365	0.0365	5.193	0.049
Residual	9	0.0633	0.00704		
Total	10	0.0999	0.00999		

Normality Test (Shapiro-Wilk) Passed (P = 0.520)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.116)

Linear Regression

Wednesday, May 22, 2024 11:28:13 AM

Data source: Data 1 in synthesis figs.JNB

$$RsaA = 5.220 - (0.0124 * Stand\ Age)$$

N = 11

R = 0.559 Rsqr = 0.312 Adj Rsqr = 0.236

Standard Error of Estimate = 0.996

	Coefficient	Std. Error	t	P
Constant	5.220	0.594	8.782	<0.001
Stand Age	-0.0124	0.00613	-2.020	0.074

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	4.046	4.046	4.082	0.074
Residual	9	8.922	0.991		
Total	10	12.968	1.297		

Normality Test (Shapiro-Wilk) Passed (P = 0.784)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.102)

Linear Regression

Thursday, May 9, 2024 11:36:43 AM

Data source: Data 1 in synthesis figs.JNB

$$\text{RshA} = 4.780 + (0.00859 * \text{Stand Age})$$

N = 11

R = 0.593 Rsqr = 0.351 Adj Rsqr = 0.279

Standard Error of Estimate = 0.632

	Coefficient	Std. Error	t	P
Constant	4.780	0.377	12.676	<0.001
Stand Age	0.00859	0.00389	2.208	0.055

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	1.945	1.945	4.875	0.055
Residual	9	3.592	0.399		
Total	10	5.537	0.554		

Normality Test (Shapiro-Wilk) Passed (P = 0.085)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.369)

Linear Regression

Tuesday, April 25, 2023 11:31:04 AM

Data source: Data 1 in synthesis figs.JNB

$$\text{RstGS} = 8.753 - (0.00691 * \text{Stand Age})$$

N = 11 Missing Observations = 1

R = 0.625 Rsqr = 0.391 Adj Rsqr = 0.324

Standard Error of Estimate = 0.466

	Coefficient	Std. Error	t	P
Constant	8.753	0.278	31.442	<0.001
Stand Age	-0.00691	0.00287	-2.405	0.040

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	1.258	1.258	5.784	0.040
Residual	9	1.957	0.217		
Total	10	3.215	0.322		

Normality Test (Shapiro-Wilk) Passed (P = 0.088)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.776)

778 **Linear Regression** Wednesday, May 22, 2024 11:29:01 AM

779

780 **Data source:** Data 1 in synthesis figs.JNB

781

782 $R_{shGS}/R_{stGS} = 0.432 + (0.00131 * Stand\ Age)$

783

784 $N = 11$

785

786 $R = 0.650$ $Rsqr = 0.422$ $Adj\ Rsqr = 0.358$

787

788 Standard Error of Estimate = 0.083

789

790 **Coefficient** **Std. Error** **t** **P**

791 Constant 0.432 0.0494 8.734 <0.001

792 Stand Age 0.00131 0.000510 2.564 0.030

793

794 Analysis of Variance:

795 **DF** **SS** **MS** **F** **P**

796 Regression 1 0.0451 0.0451 6.576 0.030

797 Residual 9 0.0617 0.00686

798 Total 10 0.107 0.0107

799

800 Normality Test (Shapiro-Wilk) Passed (P = 0.058)

801

802 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.221)

803

804 **Linear Regression** Wednesday, May 22, 2024 11:29:58 AM

805

806 **Data source:** Data 1 in synthesis figs.JNB

807

808 $R_{saGS} = 4.907 - (0.0133 * Stand\ Age)$

809

810 $N = 11$

811

812 $R = 0.647$ $Rsqr = 0.419$ $Adj\ Rsqr = 0.354$

813

814 Standard Error of Estimate = 0.848

815

816 **Coefficient** **Std. Error** **t** **P**

817 Constant 4.907 0.506 9.696 <0.001

818 Stand Age -0.0133 0.00522 -2.546 0.031

819

820 Analysis of Variance:

821 **DF** **SS** **MS** **F** **P**

822 Regression 1 4.658 4.658 6.481 0.031

823 Residual 9 6.469 0.719

824 Total 10 11.126 1.113

825

826 Normality Test (Shapiro-Wilk) Passed (P = 0.371)

827

828 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.673)

829

Wednesday, May 22, 2024 11:30:33 AM

Linear Regression**Data source:** Data 1 in synthesis figs.JNB

$$\text{RhGS} = 3.846 + (0.00639 * \text{Stand Age})$$

N = 11

R = 0.546 Rsqr = 0.298 Adj Rsqr = 0.220

Standard Error of Estimate = 0.530

	Coefficient	Std. Error	t	P
Constant	3.846	0.317	12.147	<0.001
Stand Age	0.00639	0.00327	1.955	0.082

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	1.075	1.075	3.821	0.082
Residual	9	2.531	0.281		
Total	10	3.606	0.361		

Normality Test (Shapiro-Wilk) Failed (P = <0.001)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.353)

Figure 6: Fine roots, stand age, soil respiration

Linear Regression

Wednesday, May 22, 2024 11:32:51 AM

Data source: Data 1 in synthesis figs.JNB

$$\text{Root_prod} = 1.991 - (0.00398 * \text{Stand Age})$$

N = 10 Missing Observations = 1

R = 0.426 Rsqr = 0.181 Adj Rsqr = 0.0789

Standard Error of Estimate = 0.463

	Coefficient	Std. Error	t	P
Constant	1.991	0.278	7.165	<0.001
Stand Age	-0.00398	0.00299	-1.331	0.220

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	0.380	0.380	1.771	0.220
Residual	8	1.716	0.214		
Total	9	2.096	0.233		

Normality Test (Shapiro-Wilk) Passed (P = 0.711)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.583)

Linear Regression

Wednesday, May 22, 2024 11:33:54 AM

Data source: Data 1 in synthesis figs.JNB

$$\text{RstA} = 8.191 + (0.839 * \text{Root_prod})$$

N = 10 Missing Observations = 1

R = 0.795 Rsqr = 0.632 Adj Rsqr = 0.586

Standard Error of Estimate = 0.328

	Coefficient	Std. Error	t	P
Constant	8.191	0.394	20.802	<0.001
NPPfr	0.839	0.227	3.705	0.006

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	1.476	1.476	13.728	0.006
Residual	8	0.860	0.108		
Total	9	2.336	0.260		

Normality Test (Shapiro-Wilk) Passed (P = 0.197)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.838)

917 **Burn Plots O horizon C stocks: meta-analysis**

918 **Structure: Grouped**

919 → Citation: Hedges and Olkin (1985)

920 → Effect Sizes: lnR

921 → Effect Size Variances: var_un

922 → Groups: Author

923 → Fixed Effects Model

924 → Standard confidence intervals around means based on Student's t distribution

925 → Use bootstrap for confidence intervals around means: 999 iterations

926 → Citations: Adams *et al.* (1997), Dixon (1993)

927 23 studies will be included in this analysis

928 **Group Results**

929 **Heterogeneity**

930 Source Q df P(χ^2)

931 -----

932 Gough (within) 0.49 3 0.92

933 Le Moine (within) 0.06 4 1.00

934 Lehman (within) 0.21 3 0.98

935 Nave (within) 0.02 3 1.00

936 Schaetzl (within) 1.17 5 0.95

937 Source I² 95% CI

938 -----

939 Gough (within) 0.00 0.00 to 6.30

940 Le Moine (within) 0.00 0.00 to 0.00

941 Lehman (within) 0.00 0.00 to 0.00

942 Nave (within) 0.00 0.00 to 0.00

943 Schaetzl (within) 0.00 0.00 to 0.00

944 → Citations: Higgins and Thompson (2002), Huedo-Medina *et al.* (2006)

945 Source Q df P(χ^2)

946 -----

947 Model (Between) 3.73 4 0.44

948 Error (Within) 1.94 18 1.00

949 -----

950 Total 5.67 22 1.00

951 **Mean Effect Sizes**

952 n Mean Median 95% CI Bootstrap CI Bias-corrected CI

953 -----

954 -----

955 Gough lnR 4 0.00 0.02 -1.59 to 1.59 -0.33 to 0.32 -0.40 to 0.24

956 Le Moine lnR 5 -0.04 -0.07 -1.28 to 1.20 -0.13 to 0.06 -0.13 to 0.06

957 Lehman lnR 4 -0.29 -0.33 -1.88 to 1.30 -0.51 to -0.08 -0.52 to -0.10

958 Nave lnR 4 -0.13 -0.12 -1.72 to 1.46 -0.20 to -0.07 -0.20 to -0.07

959 Schaetzl lnR 6 -1.00 -1.12 -2.05 to 0.05 -1.31 to -0.63 -1.29 to -0.61

960 -----

961 -----

962 Gough exp(lnR) 4 1.00 1.02 0.20 to 4.91 0.72 to 1.37 0.67 to 1.27

963	Le Moine exp(lnR)	5	0.96	0.93	0.28 to 3.32	0.88 to 1.06	0.88 to 1.06
964	Lehman exp(lnR)	4	0.75	0.72	0.15 to 3.67	0.60 to 0.93	0.60 to 0.90
965	Nave exp(lnR)	4	0.88	0.89	0.18 to 4.32	0.82 to 0.93	0.82 to 0.93
966	Schaetzl exp(lnR)	6	0.37	0.33	0.13 to 1.05	0.27 to 0.53	0.27 to 0.54

967 **Global Results**

968 **Heterogeneity**

969	Source	Q	df	P(χ^2)
970	-----			
971	Total	5.67	22	1.00

972 → Citation: Hedges and Olkin (1985)

973	Source	I ²	95% CI
974	-----		
975	Total	0.00	0.00 to 0.00

976 → Citations: Higgins and Thompson (2002), Huedo-Medina *et al.* (2006)

977 **Mean Effect Size**

978	n	Mean	Median	95% CI	Bootstrap CI	Bias-corrected CI	
979	-----						
980	----						
981	Global lnR	23	-0.34	-0.16	-0.78 to 0.09	-0.54 to -0.15	-0.55 to -0.16
982	-----						
983	----						
984	Global exp(lnR)	23	0.71	0.85	0.46 to 1.09	0.58 to 0.86	0.58 to 0.85

985 **Structure: Linear Regression**

986 → Citations: Hedges and Olkin (1985), Greenland (1987)

987 → Effect Sizes: lnR

988 → Effect Size Variances: var_un

989 → Independent Variable: Time

990 → Fixed Effects Model

991 → Standard confidence intervals around means based on Student's t distribution

992 → Use bootstrap for confidence intervals around means: 999 iterations

993 → Citations: Adams *et al.* (1997), Dixon (1993)

994 23 studies will be included in this analysis

995 **Regression Results**

996 **Predictors**

997	Predictor	Value	SE	95% CI	P(Normal)
998	-----				
999	Intercept	-0.48	0.47	-1.41 to 0.44	0.31
1000	Slope	0.00	0.01	-0.02 to 0.02	0.74

1001 **Heterogeneity**

1002	Source	Q	df	P(χ^2)
1003	-----			
1004	Model	0.11	1	0.74
1005	Error	5.56	21	1.00
1006	-----			
1007	Total	5.67	22	1.00

1008 **Figure 7: Stand age, soil properties, biogeochemical functions**

1009 **Linear Regression**

Monday, May 1, 2023 2:41:37 PM

1010
1011 **Data source:** Data 2 in synthesis figs.JNB

1012
1013 $\%C = 5.446 + (0.0487 * \text{Stand age})$

1014
1015 $N = 6$

1016
1017 $R = 0.851 \quad Rsqr = 0.724 \quad Adj Rsqr = 0.656$

1018
1019 Standard Error of Estimate = 1.031

1020

	Coefficient	Std. Error	t	P
1021 Constant	5.446	1.074	5.071	0.007
1022 Stand age	0.0487	0.0150	3.243	0.032

1023
1024
1025 Analysis of Variance:

1026

	DF	SS	MS	F	P
1027 Regression	1	11.181	11.181	10.514	0.032
1028 Residual	4	4.254	1.063		
1029 Total	5	15.435	3.087		

1030
1031 Normality Test (Shapiro-Wilk) Passed (P = 0.337)

1032
1033 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.060)

1034

1035 **Linear Regression**

Monday, May 1, 2023 2:42:08 PM

1036
1037 **Data source:** Data 2 in synthesis figs.JNB

1038
1039 $MBC = 295.573 + (4.579 * \text{Stand age})$

1040
1041 $N = 6$

1042
1043 $R = 0.927 \quad Rsqr = 0.859 \quad Adj Rsqr = 0.824$

1044
1045 Standard Error of Estimate = 63.617

1046

	Coefficient	Std. Error	t	P
1047 Constant	295.573	66.258	4.461	0.011
1048 Stand age	4.579	0.926	4.945	0.008

1049
1050
1051 Analysis of Variance:

1052

	DF	SS	MS	F	P
1053 Regression	1	98967.320	98967.320	24.453	0.008
1054 Residual	4	16188.680	4047.170		
1055 Total	5	115156.000	23031.200		

1056
1057 Normality Test (Shapiro-Wilk) Passed (P = 0.730)

1058
1059 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.060)

1060 **Linear Regression** Monday, May 1, 2023 2:53:15 PM

1061

1062 **Data source:** Data 2 in synthesis figs.JNB

1063

1064 $pH = 5.038 - (0.00944 * \text{Stand age})$

1065

1066 $N = 6$

1067

1068 $R = 0.861 \qquad Rsqr = 0.741 \qquad Adj \text{ Rsqr} = 0.676$

1069

1070 Standard Error of Estimate = 0.192

1071

1072 **Coefficient Std. Error t P**

1073 Constant 5.038 0.200 25.210 <0.001

1074 Stand age -0.00944 0.00279 -3.381 0.028

1075

1076 Analysis of Variance:

1077 **DF SS MS F P**

1078 Regression 1 0.421 0.421 11.434 0.028

1079 Residual 4 0.147 0.0368

1080 Total 5 0.568 0.114

1081

1082 Normality Test (Shapiro-Wilk) Passed (P = 0.752)

1083

1084 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.060)

1085

1086 **Linear Regression** Monday, May 1, 2023 3:12:45 PM

1087

1088 **Data source:** Data 2 in synthesis figs.JNB

1089

1090 $\text{AlphaG} = 7.828 - (1.320 * pH)$

1091

1092 $N = 6$

1093

1094 $R = 0.853 \qquad Rsqr = 0.728 \qquad Adj \text{ Rsqr} = 0.660$

1095

1096 Standard Error of Estimate = 0.304

1097

1098 **Coefficient Std. Error t P**

1099 Constant 7.828 1.787 4.381 0.012

1100 pH -1.320 0.404 -3.270 0.031

1101

1102 Analysis of Variance:

1103 **DF SS MS F P**

1104 Regression 1 0.990 0.990 10.692 0.031

1105 Residual 4 0.370 0.0926

1106 Total 5 1.360 0.272

1107

1108 Normality Test (Shapiro-Wilk) Passed (P = 0.775)

1109

1110 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.060)

1111

Linear Regression

Monday, May 1, 2023 3:05:30 PM

Data source: Data 2 in synthesis figs.JNB

$$\text{AlphaG} = -0.219 + (0.257 * \%C)$$

N = 6

R = 0.864 Rsqr = 0.747 Adj Rsqr = 0.684

Standard Error of Estimate = 0.293

	Coefficient	Std. Error	t	P
Constant	-0.219	0.657	-0.334	0.755
%C	0.257	0.0746	3.437	0.026

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	1.016	1.016	11.813	0.026
Residual	4	0.344	0.0860		
Total	5	1.360	0.272		

Normality Test (Shapiro-Wilk) Passed (P = 0.490)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.060)

Figure 10: Fungal 15N tracer recovery

One Way Analysis of Variance

Monday, December 11, 2023 12:41:29 PM

Data source: Data 1 in Notebook1

Dependent Variable: d15N

Normality Test (Shapiro-Wilk): Passed (P = 0.135)

Equal Variance Test (Brown-Forsythe): Passed (P = 0.821)

Group Name	N	Missing	Mean	Std Dev	SEM
Cortinariaceae	4	0	44.977	7.208	3.604
Russulaceae	7	0	31.294	5.531	2.090
Amanitaceae	3	0	33.147	6.395	3.692
Boletaceae	4	0	29.125	6.060	3.030

Source of Variation	DF	SS	MS	F	P
Between Groups	3	630.045	210.015	5.533	0.010
Residual	14	531.363	37.955		
Total	17	1161.408			

The differences in the mean values among the treatment groups are greater than would be expected by chance; there is a statistically significant difference (P = 0.010).

Power of performed test with alpha = 0.050: 0.773

All Pairwise Multiple Comparison Procedures (Fisher LSD Method):

Comparisons for factor: **Family**

Comparison	Diff of Means	LSD(alpha=0.050)	P	Diff >= LSD
Cortinariaceae vs. Boletaceae	15.852	9.343	0.003	Yes
Cortinariaceae vs. Russulaceae	13.683	8.282	0.003	Yes
Cortinariaceae vs. Amanitaceae	11.831	10.092	0.025	Yes
Amanitaceae vs. Boletaceae	4.022	10.092	0.407	No
Amanitaceae vs. Russulaceae	1.852	9.118	0.670	Do Not Test
Russulaceae vs. Boletaceae	2.169	8.282	0.583	Do Not Test

Figure 11 (Decadal change in SOC stocks)**Two Way Analysis of Variance**

Wednesday, October 27, 2021 1:51:20 PM

Data source: Data 1 in Notebook1

General Linear Model

Dependent Variable: stock_c

Normality Test (Shapiro-Wilk): Passed (P = 0.858)**Equal Variance Test (Brown-Forsythe):** Passed (P = 0.105)

Source of Variation	DF	SS	MS	F	P
year	1	38.275	38.275	6.299	0.015
horizon	2	403.051	201.526	33.163	<0.001
year x horizon	2	51.667	25.834	4.251	0.018
Residual	65	394.992	6.077		
Total	70	891.787	12.740		

Main effects cannot be properly interpreted if significant interaction is determined. This is because the size of a factor's effect depends upon the level of the other factor.

The effect of different levels of year depends on what level of horizon is present. There is a statistically significant interaction between year and horizon. (P = 0.018)

Power of performed test with alpha = 0.0500: for year : 0.621

Power of performed test with alpha = 0.0500: for horizon : 1.000

Power of performed test with alpha = 0.0500: for year x horizon : 0.600

Least square means for year :

Group	Mean	SEM
1980.000	8.415	0.417
2009.000	6.946	0.411

Least square means for horizon :

Group	Mean	SEM
A	9.233	0.503
E	4.324	0.503
B	9.486	0.514

Least square means for year x horizon :

Group	Mean	SEM
1980.000 x A	10.500	0.712
1980.000 x E	5.739	0.712
1980.000 x B	9.007	0.743
2009.000 x A	7.966	0.712
2009.000 x E	2.908	0.712
2009.000 x B	9.964	0.712

All Pairwise Multiple Comparison Procedures (Tukey Test):

1219	Comparisons for factor: horizon within 1980					
	Comparison	Diff of Means	p	q	P	P<0.050
	A vs. E	4.761	3	6.691	<0.001	Yes
	A vs. B	1.493	3	2.052	0.321	No
	B vs. E	3.268	3	4.492	0.006	Yes
1220						
1221						
1222	Comparisons for factor: horizon within 2009					
	Comparison	Diff of Means	p	q	P	P<0.050
	B vs. E	7.055	3	9.915	<0.001	Yes
	B vs. A	1.998	3	2.807	0.124	No
	A vs. E	5.058	3	7.107	<0.001	Yes
1223						
1224						
1225	Comparisons for factor: year within A					
	Comparison	Diff of Means	p	q	P	P<0.050
	1980.000 vs. 2009.000	2.534	2	3.561	0.014	Yes
1226						
1227						
1228	Comparisons for factor: year within E					
	Comparison	Diff of Means	p	q	P	P<0.050
	1980.000 vs. 2009.000	2.830	2	3.977	0.007	Yes
1229						
1230						
1231	Comparisons for factor: year within B					
	Comparison	Diff of Means	p	q	P	P<0.050
	2009.000 vs. 1980.000	0.957	2	1.315	0.356	No
1232						
1233						
1234						
1235						
1236						
1237						
1238						
1239						
1240						
1241						
1242						
1243						
1244						
1245						
1246						

Figure 12: C and water fluxes and climatology**Multiple Linear Regression**

Wednesday, May 22, 2024 12:36:27 PM

Data source: Data 1 in fluxes_thru_2020.JNB

$$\text{GPP} = -22700.350 + (11.953 * \text{Year}) - (18.329 * +\text{RC-TRT})$$

N = 34 Missing Observations = 1

R = 0.650 Rsqr = 0.422 Adj Rsqr = 0.385

Standard Error of Estimate = 79.439

	Coefficient	Std. Error	t	P	VIF
Constant	-22700.350	5175.015	-4.387	<0.001	
Year	11.953	2.575	4.643	<0.001	1.135
+RC-TRT	-18.329	29.866	-0.614	0.544	1.135

Analysis of Variance:

	DF	SS	MS	F	P
Regression	2	142991.433	71495.716	11.330	<0.001
Residual	31	195625.223	6310.491		
Total	33	338616.656	10261.111		

Column	SSIncr	SSMarg
Year	140614.569	136012.398
+RC-TRT	2376.864	2376.864

The dependent variable GPP can be predicted from a linear combination of the independent variables:

	P
Year	<0.001
+RC-TRT	0.544

Not all of the independent variables appear necessary (or the multiple linear model may be underspecified).

The following appear to account for the ability to predict GPP (P < 0.05): Year

Normality Test (Shapiro-Wilk) Passed (P = 0.440)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.131)

1300

Multiple Linear Regression

Wednesday, May 22, 2024 12:37:21 PM

1301

1302 **Data source:** Data 1 in fluxes_thru_2020.JNB

1303

1304 $ER = -13566.763 + (7.285 * Year) - (80.589 * +RC-TRT)$

1305

1306 N = 34 Missing Observations = 1

1307

1308 R = 0.555 Rsqr = 0.308 Adj Rsqr = 0.263

1309

1310 Standard Error of Estimate = 72.109

1311

	Coefficient	Std. Error	t	P	VIF
Constant	-13566.763	4697.522	-2.888	0.007	
Year	7.285	2.337	3.117	0.004	1.135
+RC-TRT	-80.589	27.110	-2.973	0.006	1.135

1316

1317 Analysis of Variance:

	DF	SS	MS	F	P
Regression	2	71782.416	35891.208	6.903	0.003
Residual	31	161190.446	5199.692		
Total	33	232972.861	7059.784		

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Column	SSIincr	SSMarg
Year	25835.295	50530.152
+RC-TRT	45947.120	45947.120

1326

1327 The dependent variable ER can be predicted from a linear combination of the independent variables:

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	P
Year	0.004
+RC-TRT	0.006

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1333 All independent variables appear to contribute to predicting ER (P < 0.05).

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1335 Normality Test (Shapiro-Wilk) Passed (P = 0.571)

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1337 Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.575)

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Multiple Linear Regression

Wednesday, May 22, 2024 12:34:30 PM

Data source: Data 1 in fluxes_thru_2020.JNB

NEE = 12729.804 - (6.464 * Year) - (79.240 * +RC-TRT)

N = 34 Missing Observations = 1

R = 0.764 Rsqr = 0.584 Adj Rsqr = 0.557

Standard Error of Estimate = 54.343

	Coefficient	Std. Error	t	P	VIF
Constant	12729.804	3540.151	3.596	0.001	
Year	-6.464	1.761	-3.670	<0.001	1.135
+RC-TRT	-79.240	20.431	-3.878	<0.001	1.135

Analysis of Variance:

	DF	SS	MS	F	P
Regression	2	128459.783	64229.891	21.750	<0.001
Residual	31	91547.223	2953.136		
Total	33	220007.005	6666.879		

Column	SSIncr	SSMarg
Year	84038.265	39775.358
+RC-TRT	44421.518	44421.518

The dependent variable NEE can be predicted from a linear combination of the independent variables:

	P
Year	<0.001
+RC-TRT	<0.001

All independent variables appear to contribute to predicting NEE (P < 0.05).

Normality Test (Shapiro-Wilk) Passed (P = 0.552)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.087)

Wednesday, May 22, 2024 12:39:59 PM

Multiple Linear Regression

Data source: Data 1 in fluxes_thru_2020.JNB

WUE = -89.783 + (0.0463 * Year) + (0.101 * +RC-TRT)

N = 34 Missing Observations = 1

R = 0.747 Rsqr = 0.558 Adj Rsqr = 0.529

Standard Error of Estimate = 0.247

	Coefficient	Std. Error	t	P	VIF
Constant	-89.783	16.616	-5.404	<0.001	
Year	0.0463	0.00826	5.597	<0.001	1.083
+RC-TRT	0.101	0.0895	1.127	0.268	1.083

Analysis of Variance:

	DF	SS	MS	F	P
Regression	2	2.383	1.191	19.543	<0.001
Residual	31	1.890	0.0610		
Total	33	4.272	0.129		

Column	SSIncr	SSMarg
Year	2.305	1.909
+RC-TRT	0.0775	0.0775

The dependent variable WUE can be predicted from a linear combination of the independent variables:

	P
Year	<0.001
+RC-TRT	0.268

Not all of the independent variables appear necessary (or the multiple linear model may be underspecified).

The following appear to account for the ability to predict WUE (P < 0.05): Year

Normality Test (Shapiro-Wilk) Passed (P = 0.302)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.661)

Linear Regression

Tuesday, May 14, 2024 9:48:58 AM

Data source: Data 1 in fluxes_thru_2020.JNB

$$\text{MAT} = -42.520 + (0.0247 * \text{Year})$$

N = 41

R = 0.309 Rsqr = 0.0957 Adj Rsqr = 0.0725

Standard Error of Estimate = 0.919

	Coefficient	Std. Error	t	P
Constant	-42.520	24.273	-1.752	0.088
Year	0.0247	0.0121	2.032	0.049

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	3.490	3.490	4.128	0.049
Residual	39	32.971	0.845		
Total	40	36.461	0.912		

Normality Test (Shapiro-Wilk) Passed (P = 0.948)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.280)

Linear Regression

Wednesday, May 22, 2024 1:16:54 PM

Data source: Data 1 in fluxes_thru_2020.JNB

$$\text{MAP} = -917.074 + (0.501 * \text{Year})$$

N = 41

R = 0.501 Rsqr = 0.251 Adj Rsqr = 0.231

Standard Error of Estimate = 10.508

	Coefficient	Std. Error	t	P
Constant	-917.074	277.388	-3.306	0.002
Year	0.501	0.139	3.612	<0.001

Analysis of Variance:

	DF	SS	MS	F	P
Regression	1	1440.310	1440.310	13.045	<0.001
Residual	39	4306.019	110.411		
Total	40	5746.329	143.658		

Normality Test (Shapiro-Wilk) Passed (P = 0.164)

Constant Variance Test (Spearman Rank Correlation): Passed (P = 0.107)

1514

Nonlinear Regression- N deposition

Wednesday, May 22, 2024 1:20:43 PM

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Data Source: Data 1 in fluxes_thru_2020.JNB

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Equation: Piecewise, 2 segment linear

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$t1 = \min(t)$

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$t2 = \max(t)$

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$region1(t) = (y1*(T1-t) + y2*(t-t1))/(T1-t1)$

1521

$region2(t) = (y2*(t2-t) + y3*(t-T1))/(t2-T1)$

1522

$f = \text{if}(t \leq T1, region1(t), region2(t))$

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R	Rsqr	Adj Rsqr	Standard Error of Estimate
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0.5096	0.2597	0.1962	0.7304
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	Coefficient	Std. Error	t	P
y1	5.1214	0.3373	15.1853	<0.0001
y2	5.0494	0.3221	15.6755	<0.0001
y3	3.8862	0.3304	11.7622	<0.0001
T1	2002.0000	7.7335	258.8753	<0.0001

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Analysis of Variance:

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	DF	SS	MS
Regression	4	898.0858	224.5215
Residual	35	18.6721	0.5335
Total	39	916.7579	23.5066

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Corrected for the mean of the observations:

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	DF	SS	MS
Regression	3	6.5489	2.1830
Residual	35	18.6721	0.5335
Total	38	25.2210	0.6637

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Statistical Tests:

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Normality Test (Shapiro-Wilk) Passed (P = 0.3501)

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W Statistic= 0.9690 Significance Level = 0.0500

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Constant Variance Test (Spearman Rank Correlation) Passed (P = 0.7571)

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Soil respiration: US-UMB vs. US-UMd

Factor	df	t	P
Footprint	10	14.781	0.00000004

Group Means

UMd	6.54
UMB	7.37

Difference of Means

0.83

For Review Only

Appendix S1 References

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- 1636

Author	Control_age	Stand_age	CTRL C stock	TRT C stock	lnR	var_un
Schaetzl 1994	90	12	2	0.5	-1.492	1
Schaetzl 1994	90	38	2	0.5	-1.328	1
Schaetzl 1994	90	44	2	0.7	-1.064	1
Schaetzl 1994	90	56	2	0.9	-0.832	1
Schaetzl 1994	90	69	2	1.8	-0.128	1
Schaetzl 1994	90	81	2	0.6	-1.171	1
Nave 2019	103	16	4.4	3.8	-0.157	1
Nave 2019	103	34	4.4	4.1	-0.064	1
Nave 2019	103	60	4.4	4.1	-0.077	1
Nave 2019	103	78	4.4	3.6	-0.216	1
Gough 2007	68	6	6.2	3.8	-0.495	1
Gough 2007	68	24	6.2	7.2	0.149	1
Gough 2007	68	50	6.2	9.8	0.459	1
Gough 2007	68	56	6.2	5.6	-0.11	1
Lehman 1993	82	57	16.6	17	0.024	1
Lehman 1993	82	45	20.1	16.9	-0.175	1
Lehman 1993	82	39	18.4	11.3	-0.486	1
Lehman 1993	82	18	22.1	13.1	-0.526	1
Le Moine 2012	101	76	8	7.7	-0.029	1
Le Moine 2012	101	64	8	7.4	-0.07	1
Le Moine 2012	101	58	8	7.4	-0.07	1
Le Moine 2012	101	32	8	9.2	0.147	1
Le Moine 2012	101	14	8	6.6	-0.183	1

Author identifies the source dataset

Control_age indicates which stand age was used as the control for each source dataset

Stand_age indicates the stand age, in years, since the experimental cut+burn manipulation

CTRL C stock is the O horizon C stock in Mg ha-1 for the control condition

TRT C stock is the O horizon C stock in Mg ha-1 for the treatment plot

lnR is the ln-transformed response ratio

var_un is set at 1 in order to conduct an unweighted meta-analysis (all weights the same)