Tree diversity increases carbon stocks and fluxes above- but not belowground in a tropical forest experiment

Running title: Tree diversity increases carbon sequestration

Florian Schnabel[1,2,3,*], Joannès Guillemot[4,5,6], Kathryn E. Barry[7], Melanie Brunn[8,9], Simone Cesarz[3,10], Nico Eisenhauer[3,10], Tobias Gebauer[11,12], Nathaly R. Guerrero-Ramirez[13,14,15,16], I. Tanya Handa[13], Chris Madsen[17], Lady Mancilla[18], Jose Monteza[18], Tim Moore[19], Yvonne Oelmann[20], Michael Scherer-Lorenzen[11], Luitgard Schwendenmann[21], Audrey Wagner[22,23], Christian Wirth[2,3,24], Catherine Potvin[17,18]

[1]Chair of Silviculture, Institute of Forest Sciences, University of Freiburg, Freiburg, Germany

[2]Systematic Botany and Functional Biodiversity, Leipzig University, Leipzig, Germany

[3]German Centre for Integrative Biodiversity Research (iDiv) Halle-Jena-Leipzig, Leipzig, Germany

[4]CIRAD, UMR Eco&Sols, Montpellier, France

[5]Eco&Sols, Univ. Montpellier, CIRAD, INRAe, Institut Agro, IRD, Montpellier, France

[6]Department of Forest Sciences, ESALQ, University of São Paulo, Piracicaba, São Paulo, Brazil

[7]Ecology and Biodiversity, Department of Biology, Institute of Environmental Biology, Utrecht University, Utrecht, Netherlands

[8]Institute for Integrated Natural Sciences, University of Koblenz, Koblenz, Germany

[9]Institute for Environmental Sciences, RPTU University of Kaiserslautern-Landau, Landau, Germany

[10]Institute of Biology, Leipzig University, Leipzig, Germany

[11]Gebotany, Faculty of Biology, University of Freiburg, Freiburg, Germany

[12]Geo-konzept society of environmental planning GmbH, Adelschlag, Germany

[13]Département des sciences biologiques, Université du Québec à Montréal, Montréal, Canada

[14]Biodiversity, Macroecology and Biogeography, Faculty of Forest Sciences and Forest Ecology, University of Göttingen, Göttingen, Germany

[15]Silviculture and Forest Ecology of Temperate Zones, Faculty of Forest Sciences and Forest Ecology,

University of Göttingen, Göttingen, Germany

[16]Centre of Biodiversity and Sustainable Land Use, University of Göttingen, Göttingen, Germany

[17] Neotropical Ecology Laboratory, Department of Biology, McGill University, Montreal, QC, Canada

[18]Smithsonian Tropical Research Institute, Panama City, Panama

[19]Department of Geography, McGill University, Montreal, QC, Canada

[20]Geoecology, Department of Geosciences, Tübingen University, Tübingen, Germany

[21]School of Environment, University of Auckland, Auckland, New Zealand

[22]Faculty of Agricultural and Environmental Sciences, McGill University, Montreal, QC, Canada

[23]Nature-based Solutions Initiative, Department of Biology, University of Oxford, Oxford, United Kingdom

[24]Max Planck Institute for Biogeochemistry, Jena, Germany

*Corresponding author. E-mail: florian.schnabel@waldbau.uni-freiburg.de, Tel: +49/761/2033679

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Abstract

International commitments advocate large-scale forest restoration as a nature-based solution to climate change mitigation through carbon (C) sequestration. Mounting evidence suggests that mixed compared to monospecific planted forests may store more C, exhibit lower susceptibility to climate extremes and offer a broader range of ecosystem services. However, experimental studies comprehensively examining the control of tree diversity on multiple C stocks and fluxes above- and belowground are lacking. To address this gap, we leverage data from the Sardinilla experiment in Panama, the oldest tropical tree diversity experiment which features a gradient of one-, two-, three- and five-species mixtures. Over 16 years, we measured multiple above- and belowground C stocks and fluxes, ranging from tree aboveground C, over leaf litter C production, to soil organic carbon (SOC). We show that tree diversity significantly increased aboveground C stocks and fluxes, with a 57% higher gain in aboveground tree C in five-species mixtures compared to monocultures (35.7 ± 1.8 *vs* 22.8 ± 3.4 Mg C ha^{-1}) 16 years after planting. In contrast, we observed a net reduction in SOC (on average -11.2 ± 1.1 Mg C ha^{-1}) and no significant difference in SOC$_3$ stocks (the tree-derived SOC fraction) between five-species mixtures and monocultures (13.0 ± 0.9 *vs* 15.1 ± 1.3 Mg C ha^{-1}). Positive tree diversity effects persisted despite repeated climate extremes and strengthened over time for aboveground tree growth. Higher tree growth in mixtures enhanced leaf litter and coarse woody debris C fluxes to the soil, resulting in a tightly linked C cycle aboveground. However, the only link between above- and belowground C stocks and fluxes was between leaf litter C inputs and SOC$_3$ which was positively affected by diversity. Our study elucidates the mechanisms through which higher tree diversity bolsters the climate mitigation potential of tropical forest restoration. Restoration schemes should prioritize mixed over monospecific planted forests.

Introduction

Forest restoration is promoted as a key strategy for mitigating climate change through carbon (C) sequestration. Several global initiatives, such as the Bonn Challenge and the New York Declaration on forests, aim to restore 350 Mha of forests by 2030. Forest restoration, in particular in the tropics, has the most significant climate mitigation potential of 20 proposed nature-based solutions, potentially sequestering up to 10.1 PgCO$_2$ equivalents per year (Griscom et al., 2017). However, massive reforestation efforts should ensure the protection of land for agriculture (Dooley et al., 2022) and avoid the replacement of other ecosystems, such as natural grasslands (Parr et al., 2024; Seddon, 2022). A solution is to target the vast areas of degraded land suited for forest growth (Bauhus et al., 2010). Current reforestation pledges largely focus on monospecific planted forests (Lewis et al., 2019) despite the mounting evidence that tree species-diverse planted forests (hereafter mixed planted forests or mixtures) can exhibit lower susceptibility to stress and disturbances such as droughts and storms while simultaneously providing a broader range of ecosystem services such as C sequestration, biodiversity conservation and cultural services at higher levels than monospecific plantations (Messier et al., 2021). Consequently, mixed planted forests better fulfil current international targets such as the Kunming-Montréal Global Biodiversity Framework (CBD, 2022).

Biodiversity-ecosystem function theory suggests that mixed planted forests may outperform monocultures in terms of productivity, through complementary resource partitioning across species, abiotic facilitation or biotic feedbacks (Barry et al., 2019). Thus, mixed planted forests may also outperform monocultures with respect to their role in climate regulation. Indeed, there is accumulating evidence that mixed planted forests can store more C than their monoculture counterparts (Chen et al., 2023; Lecina-Diaz et al., 2018; Messier et al., 2021; van der Sande et al., 2017; Warner et al., 2023; Xu et al., 2020). More importantly, mixtures may also be more stable than monocultures in the face of climate extremes or climate variability in general (Isbell et al., 2018), as some species may 'insure' the community against the reduced functioning of other species (Yachi and Loreau, 1999). Mixed planted forests indeed feature higher temporal stability of biomass

production than monocultures during periods with variable climatic conditions, including particularly wet

and dry years (Jucker et al., 2014; Schnabel et al., 2021). However, most existing studies assessed (or

indirectly inferred) tree diversity effects on C stocks, fluxes, and stability in terms of aboveground tree C

(AGC), with fewer studies examining root C or soil organic C (SOC) (e.g. Xu et al., 2020) and even fewer

ones the multiple C fluxes above- and belowground, connecting these C pools.

We posit that tree species richness (hereafter tree diversity) may affect C stocks and fluxes both above- and

belowground. C stocks refer to the C stored in reservoirs such as AGC or SOC, whereas C fluxes are the

flow of C between these reservoirs over time. We anticipate comparable diversity effects above- and

belowground as has been shown in grassland experiments (Ravenek et al., 2014; Weisser et al., 2017).

Indeed, there is evidence for significant tree diversity effects on various C fluxes, ranging from enhanced

leaf litter (Huang et al., 2017) and coarse woody debris (CWD) production (Liu et al., 2018) to enhanced

microbial respiration and thus decomposition (Chen et al., 2020). Ultimately, the balance of these different

C fluxes determines net tree diversity effects on C sequestration in forests (Liu et al., 2018). For example,

tree diversity is often reported to increase tree biomass production (hence C gain, Potvin et al., 2008) but

may, in some cases, also increase tree mortality (hence C loss, Searle et al., 2022). Similarly, tree diversity

may increase SOC through diversity-induced enhancements of C inputs into the soil via plant litter or root

exudate production, but changes in soil community and functioning may also enhance C losses due to

decomposition (Chen et al., 2020; Handa et al., 2014; Lange et al., 2015). As a result no or no coherent

effect of tree diversity was also reported (e.g. Martin-Guay et al., 2022; Scherer-Lorenzen et al., 2007).

Comprehensive assessments of the multiple C stocks and fluxes in forests and their intricate relationships

are scarce (Xu et al., 2020). A notable exception is Liu et al. (2018), who studied naturally established

subtropical forests in China, revealing significant positive effects of tree diversity on AGC, root C, CWD,

and SOC, and significant correlations between tree diversity and AGC, CWD, and leaf litter production.

However, in complex natural environments like forests, environmental variation and tree diversity

interactively influence carbon stocks and fluxes (van der Sande et al., 2017). Despite attempts using

structural equation models (SEMs) to identify direct and indirect relationships (Chen et al., 2018; Chen et

al., 2023; Li et al., 2020), mechanistically disentangling these drivers remains challenging in observational

studies. Planted tree diversity experiments, which were specifically designed to compare monocultures and

mixtures of increasing diversity while controlling for environmental variation (Depauw et al., 2024;

Scherer-Lorenzen et al., 2005), offer an ideal setting for elucidating linkages among C stocks and fluxes.

Until recently, the young age of most tree diversity experiments, the slow development of trees in boreal

and temperate experiments and the different response times and dynamics of C compartments over the

course of stand development (e.g. faster responses of aboveground compared to belowground C; Ravenek

et al., 2014) prevented analysing the temporal dynamics of tree diversity effects on C stocks and fluxes.

Moreover, a temporal perspective on C residence time, i.e. the time C is stored within a reservoir, is a

prerequisite for assessing the stability of C storage under climate variability and investigating whether tree

diversity's control on C stocks and fluxes increases as forest stands develop. Increases in ecosystem

functioning over time in more diverse tree communities have been demonstrated for aboveground tree

productivity (Guerrero-Ramírez et al., 2017; Jucker et al., 2020) but not for multiple C stocks and fluxes

and their relationships.

Here, we use data on temporal changes in ten C-related stocks and fluxes measured in the oldest tropical

tree diversity experiment, the Sardinilla experiment established in 2001 in Panama, which is part of the

global network of tree diversity experiments (TreeDivNet). After two decades of C-related research (Cesarz

et al., 2022; Coll et al., 2008; Guerrero-Ramírez et al., 2016; Guillemot et al., 2020; Hutchison et al., 2018;

Kunert et al., 2019; Kunert et al., 2022; Madsen et al., 2020; Moore et al., 2018; Murphy et al., 2008; Potvin

et al., 2011; Ruiz-Jaen and Potvin, 2011; Sapijanskas et al., 2013; Sapijanskas et al., 2014; Scherer-

Lorenzen et al., 2007; Schnabel et al., 2019; Wolf et al., 2011) and due to the comparably fast tree growth

in the tropics, the Sardinilla experiment features a wealth of C-related variables above- and belowground

for planted forests with 1–5 tree species and considerably large-sized trees (with the tallest trees over 25

m), which we leverage here to explore tree diversity effects on C stocks and fluxes across 16 years (2001–

2017). Since planting, the experiment experienced repeated climate extremes including a severe El Niño-

driven drought and a Hurricane (see methods). Although it is not possible to disentangle the intertwined

impacts of stand development and these climate extremes, their occurrence has provided us with the unique

opportunity to evaluate the role of tree diversity for C stocks and fluxes in the face of severe climate events.

We anticipate a stronger positive tree diversity effect at later stages of stand development, due to enhanced

ecosystem functioning in more diverse tree communities over time (Guerrero-Ramírez et al., 2017) and a

higher stability of diverse communities to climatic extremes (Schnabel et al., 2021). Specifically, we tested

the following hypotheses: (H1) C stocks and fluxes increase with increasing tree diversity. (H2) Positive

tree diversity effects on C stocks and fluxes increase with stand development despite repeated climate

extremes. Finally, we use SEMs to test how C stocks, fluxes, and their control through tree diversity are

connected through direct and indirect relationships above- and belowground using ten explicit hypotheses

(Supplementary Table S1).

Methods

Description of the study site

This study is based on data collected over 16 years in the Sardinilla planted forest. Established in 2001

(Scherer-Lorenzen et al., 2005) Sardinilla is the oldest experiment of the International Network of Tree

Diversity Experiments (TreeDivNet; https://treedivnet.ugent.be/). The site was originally planted with six

native tree species, namely *Luehea seemannii* Triana & Planch (Ls), *Cordia alliodora* (Ruiz & Pavon) Oken

(Ca), *Anacardium excelsum* (Bert. & Balb. Ex Kunth) Skeels (Ae), *Hura crepitans* L. (Hc), *Tabebuia rosea*

(Bertol.) DC. (Tr) and *Cedrela odorata* L. (Co). A total of 24 plots ranging from 0.2025 to 0.2304 ha

(approximately 45 x 45 m) were established following a randomized complete block design. Of these, 12

plots were monocultures (2 plots per species), six plots contained a three-species mix with each species

present in two plots, and six plots contained all tree species. Due to high mortality experienced by Ca in the

two years after planting, only 22 plots were maintained over the 16 years of the experiment. This paper thus

considers the effect of three diversity levels, grouped as 1, 2-3, and 5 species. Elevation across the site

ranges from a ridge at 79 m ASL to low areas at 67 m ASL (Healy et al., 2008) resulting in a gradient of

soil types ranging from Vertic Luvisol on the ridge to Gleyic Luvisol in the low part of the plantation

(Oelmann et al., 2010). Despite the low pH, soils of the Sardinilla plantation are characterized by high

nutrient availability because of the carbonate-rich parent material (Oelmann et al., 2010). In the first 10

years of plantation establishment, soil organic matter decreased by 18% in the plantation (Moore et al.,

2018). Further details on the Sardinilla tree diversity experiment can be found in Scherer-Lorenzen et al.

(2005) and Potvin and Dutilleul (2009).

We examined three periods: an early (p1), a mid (p2), and a late period (p3) of plantation development (Fig.

S1). These periods were characterized by repeated climate extremes, with the mid-period featuring an

extremely wet year (2010) and the late period a severe El Niño-driven hotter drought (2015) triggering

growth reductions and elevated tree mortality (Browne et al., 2021; Detto et al., 2018; Hutchison et al.,

2018; Schnabel et al., 2019). Subsequently, in November 2016, the experiment was hit by Hurricane Otto,

a tropical storm that formed off the coast of Panama in the Caribbean Sea inducing stem breakages in the

experiment. Climatic conditions at the Sardinilla experiment were characterized in terms of annual mean

temperature, precipitation sum, and drought index (Standardized Precipitation Evapotranspiration Index

(SPEI; Vicente-Serrano et al., 2010)), with all climate variables illustrating the climate extremes described

above in terms of temperature and precipitation extremes and drought conditions (Supplementary Fig. S2).

Data collection

We measured ten compartments of the forest C cycle, namely aboveground tree C (AGC), tree coarse root

C (CRC), coarse woody debris C (CWDC), C in herbaceous biomass (herbaceousC), leaf litter C production

(litterC), soil organic C (SOC), leaf litter decomposition, root decomposition, soil microbial biomass C

(C_{mic}), and soil respiration. In addition, we included canopy opening as a co-variable, with potentially

important influences on C stocks and fluxes. Each variable was measured during three periods of plantation

development: an early (2001 for SOC and 2005, 2006 & 2007 for the other variables), a mid (2011, 2012

& 2013), and a late period (2016 & 2017) (see Supplementary Fig. S1 for a timeline). Three variables were

only measured in some periods: C_{mic} in the mid (2013) and late (2017), root decomposition in the mid

(2012), and soil respiration in the late (2017) period. If not stated otherwise, the sample size for all variables

was n = 22 plots. We aggregated variables to periods in our analysis as not all variables were measured in

all years. To scale individual tree measurements up to community measures, we used diameter and height

inventories of all trees in the plantation conducted at the end of each growing season (December-January)

in 2005, 2012, and 2016. The measurement of individual variables is described briefly below, with details

provided in the Supplementary Methods.

Aboveground tree carbon

Aboveground tree biomass (AGB) estimates were based on species- and diversity-specific allometric

equations developed after harvesting and measuring 150 and 167 trees in the experiment in 2005 and 2017.

AGB was calculated as the sum of trunk and branch biomass (excluding leaves to focus on the more

permanent C-components of the trees). Allometric models are provided in Supplementary Material. The

best-fitting models were then combined with annual diameter and height inventories of all trees in the

experiment conducted at the end of each growing season (December-January) to estimate the AGB of each

tree in each period. Allometric models calibrated in 2017 were used for the mid and late period. AGB was

converted to aboveground tree C (AGC) using species-specific trunk C concentrations (Elias and Potvin,

2003).

Coarse root carbon

To estimate coarse root C (CRC) we relied on root:shoot ratios from the Sardinilla site based on two

different root excavation campaigns. For CRC in the early period, we relied on root:shoot ratios developed

in 2004 from excavating of three-year-old trees (Coll et al., 2008). For CRC values in the mid and late

period, we used species-specific root:shoot ratios developed in 2017 (Guillemot et al., 2020). The species-

specific root:shoot ratios were then multiplied with AGC to obtain CRC estimates of all trees in the

experiment.

Coarse woody debris carbon

All visible branches and stems fallen on the ground were collected annually in each plot and weighted to

obtain a measure of coarse woody debris (CWD) biomass. We used samples collected at the end of the dry

season (around March), except for the early period, where the collection lasted from July to August. CWD

was put back after weighing. CWD biomass was converted to C (hereafter CWDC) using the species-

specific trunk C concentration detailed above.

Herbaceous carbon

Herbaceous vegetation was cut, dried, and weighed in four quadrats (0.5 m^2) per plot, see Potvin et al.

(2011). C concentration was determined using an elemental analyser, and herbaceous biomass was

converted to herbaceous C using the average C concentration (42.72%) of legumes and grasses/non-

leguminous herbs.

Leaf litter carbon

Leaf litter was collected bi-weekly in 3–6 litter traps of 1 m^2 per plot, with trap numbers varying with the

number of tree species per plot. Each trap was positioned 1 m away from a tree of each species present in

each plot, see Scherer-Lorenzen et al. (2005). The litter was dried at a constant temperature of 70°C. Leaf

litter production was calculated by dividing total dry biomass in grams from each trap by the number of

days between two litter collection dates to determine the rate of litter fall per day per m^2. Litter biomass

production was converted to litter C production (hereafter 'litter C') using plot- and species-specific carbon

concentrations from dry season litter (Scherer-Lorenzen et al., 2007).

Soil organic carbon

Four soil cores were collected from each plot during plantation establishment (2001) and in the mid and

late periods, dried, and analysed for bulk density, SOC concentration (%), and $\delta^{13}C$ values. SOC (kg m^{-2})

stock was calculated from bulk density and C concentration. We examined not only SOC but also its C_3-

and C_4-derived fractions since the latter is associated with the C_4 grasses within the herbaceous vegetation

and in the pasture that existed prior to the plantation establishment, while the former is associated with the

C_3 inputs via litter of the trees and C_3 herbaceous plants (Moore et al., 2018). Assuming a C_3 plant $\delta^{13}C$

input of -28 ‰ and a residual C_4 plant $\delta^{13}C$ of -13 ‰, estimates of the percentage and mass of C_3-plant

derived SOC (SOC_3) and C_4-plant derived SOC (SOC_4) were made (Moore et al., 2018), where SOC_3 and

SOC_4 are percentages of total SOC that add up to one hundred percent. This approach allowed us to

determine the temporal changes in SOC derived predominately from trees (SOC_3) and C_4 grasses in the

former pasture (SOC_4).

Leaf litter decomposition

Leaf litter decomposition, hereafter litter decomposition, was measured using nylon bags filled with dry

litter from litter traps in the center of the plots. For species mixtures, equal proportions of litter from each

species were used, see Scherer-Lorenzen et al. (2007). Litter decomposition was measured in n = 17 plots

in the early and mid and n = 11 plots in the late period. Mass loss was determined by drying and weighing

the remaining litter. The percent mass remaining was divided by the number of days of decomposition for

standardization and averaged per plot.

Root decomposition

274 Root decomposition was measured using the root material of the five tree species using nylon mesh bags

275 filled with dry roots of 4th and 5th orders (Guerrero-Ramírez et al., 2016). Decomposition bags were installed

276 in the ten monocultures and three five-species mixture plots (n = 13). Mass loss was determined by washing

277 and drying the roots and, as for litter decomposition, the percent mass remaining was divided by the number

278 of days of decomposition.

Soil microbial biomass

281 C_{mic} was measured as substrate-induced respiration, that is, the respiratory response of microorganisms to

282 glucose addition (Anderson and Domsch, 1978). C_{mic} was calculated according to Beck et al. (1997) using

283 the same analytical devices (RMS Schuller, Darmstadt, Germany) for all measurements, see Cesarz et al.

284 (2022).

Soil respiration

287 Soil respiration was measured with a portable infrared gas analyser equipped with a soil respiration chamber

288 at six to eight randomly chosen locations per plot in the wet season (late July and early August). Changes

289 in CO_2 concentration over time were recorded every 5 sec over 90–120 sec when pressing the chamber

290 gently on the forest.

Canopy opening

Canopy opening was measured using four hemispheric photos per plot in October or November when trees

were fully leaved-out. Photos were analysed by the Gap Light Analyser (GLA) program (Frazer et al., 1999)

to extract canopy structure from hemispheric photos, see Sapijanskas et al. (2014). Canopy opening (a gap

fraction in %) can be considered a measure of canopy space-filling, which may mediate tree diversity effects

on C stocks and fluxes (see Supplementary Table S1).

Data Analysis

We expressed the stock and flux variables in a common unit of 1 m^2 to avoid extrapolating variables that

were measured only in small areas. We considered AGC, CRC, SOC and C_{mic} as stock variables. AGC and

CRC (kg) of individual trees were summed per plot and then expressed at the scale of one m^2 to ensure

comparability of all measurements and to account for slight variations in plot size (plots ranging from

0.2025 to 0.2304 ha). For SOC (initially measured as kg m^{-2}), the data were averaged at the plot level for

each year. We also calculated changes in stocks between the three observation periods as:

$\Delta stock = \frac{(stock_{t+1} - stock_t)}{time}$ (1)

where *stock* is either AGC, CRC or SOC and its fractions SOC_3 and SOC_4 in period *t* and *time* is the number

of years and months between two measurements (see Supplementary Fig. S1 for a timeline) resulting in

Δstock estimates in kg C m^{-2} year^{-1}. We considered CWDC, herbaceousC, litterC, leaf litter decomposition,

root decomposition, and soil respiration as annual flux variables. CWDC produced in one year was found

to mostly decompose until the end of the wet season in each year of our observation period. Similarly,

herbaceousC regrew each year and dead herbaceous material decomposed completely within a year. We

therefore considered annual measurements of CWDC and herbaceousC in our tropical forest system as

annual fluxes (kg C m^{-2} year^{-1}) rather than as stocks, as this attribution more closely reflected the reality in

the examined tropical forest compared to calculating changes in these variables between several years. The

C flux variables were analysed as follows. CWDC was estimated at the level of the plots and down-scaled

to 1 m^2 accounting for plot size. HerbaceousC was averaged at the plot level for each year. LitterC (kg C

m^2 day^{-1}) was averaged per plot and across the different collection dates and then scaled to an annual flux

(kg C m^2 year^{-1}). Litter and root decomposition was averaged across the different samples within one plot

and expressed in % mass remaining; decomposition data was not analysed jointly with other variables as

decomposition was not measured for all plots in all periods. Microbial biomass (μg C$_{mic}$ g^{-1} dry weight soil)

and soil respiration (μmol m^{-2} s^{-1}) were calculated as average values across measurement locations per plot.

Canopy opening, our co-variable, was averaged across the different samples within one plot and expressed

in %.

Multivariate Analyses of Variance

A snapshot of the compartments of the forest C cycle after 16 years of growth (2016-2017) was obtained

by Multivariate Analyses of Variance (MANOVA) testing the effect of tree diversity for different

compartments expected to be correlated. Two MANOVAs examined the tree biomass related variables

(AGC, CRC and CWDC) and the soil related ones (SOC, SOC$_4$ and SOC$_3$). A third MANOVA analysed

the effect of tree diversity on canopy opening and litterC considering that both variables might be correlated.

The fourth MANOVA considered soil microbial biomass and soil respiration. The analyses were performed

using Proc GLM of SAS version 9.4.

Understanding the build-up of the diversity effects across the examined periods proceeded using mixed-

effects Analyses of Variance (mixed-effects ANOVAs). As we were interested in temporal dynamics, we

focused on annual C fluxes and changes in stocks rather than on stocks per se to ensure a better

comparability of the examined variables and to avoid legacy effects present in stock variables (see e.g.

Chen et al. (2023)). The mixed-effects ANOVAs were performed for each Δstock and flux variable at the

plot level, according to the following model:

$R = \beta0 + \beta1 \times \text{diversity} + \beta2 \times \text{time} + \beta3 \times (\text{diversity} \times \text{time}) + b_i + \varepsilon_{ij}$ (2)

where R is the respective response variable and *diversity* had four levels corresponding to the number of

species planted per plot (1, 2, 3, and 5), and *time* had three levels (early, mid, and late period). β_0, β_1 and

β_3 are the fixed effect coefficients, b_i is the random effect for experimental plot accounting for repeated

measurements and ε_{ij} the error term. The model assumes the random effect b_i to be normally distributed

with mean and variance of $N(0, \sigma^2)$. Our mixed-effects ANOVAs allow testing for the presence of a time

by diversity effect, namely a differential build-up of C stocks and fluxes through time in response to tree

diversity. ANOVAs were used to test for diversity effects on root decomposition and soil respiration which

were only measured once. Extreme values and model assumptions, including normality and

heteroscedasticity, were checked visually and with Shapiro-Wilk test (see also Supplementary Analysis 1).

We decided to only remove two CWDC data points, as in all other cases, a plausible biological explanation

existed. Mixed-effects ANOVAs were fit in R version 4.3.0 with the packages lme4 (Bates et al., 2015) and

lmerTest (Kuznetsova et al., 2017). Least-squares means were estimated with the emmeans package (Lenth,

2020).

Structural equation models

To understand the mechanisms underlying tree diversities control on C dynamics across time, we used

structural equation models (SEMs). We developed a hypothesis-driven conceptual model based on *a priori*

knowledge of mechanisms that may drive and relate C stocks and fluxes in forest ecosystems (Fig. S3;

Table S1). This approach enabled us to test the direct and indirect relationships between C stocks, fluxes,

and tree diversity. Indirect relationships are those that are mediated by other variables. We tested whether

tree diversity affected C stocks and fluxes indirectly via diversity-induced decreases in canopy opening due

to enhanced canopy space-filling or increases in tree growth through examining relationships between tree

species richness, canopy opening, and tree growth (expressed as ΔAGC). Moreover, as we assumed canopy

opening and ΔAGC to be correlated, we included a partial correlation between both. We subsequently

expected canopy opening and tree growth to influence herbaceousC, litterC and CWDC in that (1)

decreased canopy at high diversity would correlate negatively with herbaceousC but positively with litterC

while (2) enhanced tree growth at high diversity would correlate positively with litterC and CWDC. We

did not include direct pathways between tree diversity and these variables as we expected tree diversity

effects to be predominately mediated by canopy opening or tree growth. Finally, we expected herbaceousC,

litterC, and CWDC to be the main aboveground C inputs to the soil, hypothesizing that they would

positively influence and correlate with ΔSOC and ΔCRC. We used the summed belowground ΔC stock

($\Delta SOC_3 + \Delta CRC$) to test this assumption to avoid the need to fit too many pathways considering sample

size (n = 22 plots). As we were interested in tree diversity effects, we focused on SOC_3, the tree-derived

fraction of SOC, but also tested the same SEMs for SOC (sum of SOC_3 and SOC_4). Finally, we tested for

potential direct effects of tree diversity on belowground ΔC stock not mediated by the tested relationships.

To examine temporal trends, we fit separate SEMs for the three periods of plantation development. As

information on ΔSOC, a crucial variable for our SEMs, was only available for the mid and late period (see

Fig. S1) we fit SEMs only for these two periods.

All SEMs focussed on C fluxes and changes in stocks to ensure a better comparability between variables.

Moreover, we only included variables available in kg C m^2 year^{-1}, except for canopy opening. We used

piecewise SEMs (Lefcheck, 2016) to test the relative importance of and support for these hypothesized

pathways. Global model fit was assessed via Fisher's C statistic (P>0.05). We assessed the independence

of variables and included partial, non-directional correlations to improve model fit based on tests of directed

separations (P<0.05 for violation of independence claims). For each SEM we calculated standardized path

coefficients, scaled by the standard deviations of the variables, which allowed us to compare the strength

of paths within and among models (Lefcheck et al., 2018). Individual pathways were fit as linear models

considering the number of species planted per plot (1, 2, 3, and 5) as continuous and not as categorical

variable as in the mixed-effects ANOVAs. SEMs were fit with the package piecewiseSEM (Lefcheck,

2016) and linear mixed-effects models with the packages lme4 (Bates et al., 2015) and lmerTest

(Kuznetsova et al., 2017) in R version 4.3.0.

Results

C stocks and fluxes after 16 years of tree growth: a snapshot in time

Over a period of 16 years, the experimental tree plantation accumulated an average of 35.9 ± 2.7 Mg C ha^{-1} in the trees (AGC + CRC), while SOC decreased on average by 11.2 ± 1.1 Mg C ha^{-1}, resulting in a net gain of 24.7 ± 2.9 Mg C ha^{-1} or 90.7 ± 10.6 Mg CO_{2eq} ha^{-1}. MANOVA unveiled a significant effect of tree diversity on the C compartments directly related to trees: AGC, CRC and CWDC (Table 1, Fig. 1). The diversity effect was mainly driven by AGC (Fig. 1) with both CRC and CWDC being significantly correlated to AGC (0.708, p=0.0007 and 0.624, p=0.004, respectively). In 2017, according to Tukey Studentized range test, tree AGC in the 5-species plot, was, with 35.7 ± 1.8 Mg C ha^{-1}, significantly higher than in monocultures (22.8 ± 3.4 Mg C ha^{-1}), a 57% increase (Fig. 1). The MANOVA computed with the canopy-related variables (canopy opening and litterC) also showed a significant effect of diversity (Table 1) with a significant negative correlation between the two variables (-0.741, p=0.0004). This diversity effect was predominantly driven by litterC that was 64% higher in 5-species mixtures than in monocultures. At $1.1–1.4$ Mg C ha^{-1} (Fig. 1; Fig. S4) herbaceousC played a role similar to litterC in the system, albeit it did not respond significantly to diversity (Table 2). None of the two MANOVAs performed on soil-related C compartments detected a significant effect of tree diversity (Table 1).

Fig. 1 C stocks and fluxes after 16 years of tree growth. Shown are means and standard errors of the C stocks in Mg
C ha^{-1} (brown boxes) and fluxes in Mg C ha^{-1} year^{-1} (blue boxes); numbers printed in bold. Variables in other units,
including canopy opening and litter decomposition given in (%), soil respiration given in µmol m^{-2} s^{-1} and microbial
biomass given in µg C$_{mic}$ g soil dw^{-1} are not printed in bold to allow for separation. The size of the boxes in the 5-
species mixture are scaled relative to the monoculture, with diversity-induced increases or decreases in C stocks and
fluxes indicated by larger or smaller boxes, respectively. An overview of all analysed mixtures (2-, 3- and 5-species
mixtures) is shown in Fig. S4. The sum of SOC$_3$ and SOC$_4$ gives SOC. AGC: aboveground tree C, CRC: coarse root
C, CWDC: coarse woody debris C, SOC$_4$: C$_4$ derived soil organic C (SOC), SOC$_3$: C$_3$ derived SOC.

Table 1 MANOVAs for C stocks and fluxes after 16 years of growth. The main effect tested was tree diversity with 4
different levels (1, 2, 3 and 5 species per plot). Abbreviations are given in Fig. 1. Significant effects (p <0.05) printed
in bold.

Compartments	Roy's Greatest Root	$F_{3,18}$	p-value
AGC/CRC/CWDC	0.8975	5.39	**0.0080**
LitterC/Canopy opening	0.5974	3.58	**0.0343**

SOC/SOC$_4$/SOC$_3$	0.2059	1.24	0.3260
Microbial biomass/Soil respiration	0.1722	1.03	0.4015

Diversity effects on changes in C stocks and fluxes across time

With a least-squares mean of 0.140 kg C m^2 yr^{-1}, tree ΔAGC increment was significantly slower in

monocultures than in most mixtures, which had increments of 0.140, 0.232, and 0.226 kg C m^2 yr^{-1} in 2-, 3-

, and 5-species mixtures, respectively (Table 2, Fig. 2). The positive tree diversity effect on ΔAGC tended

to strengthen over time in the 5-species mixtures (p = 0.16 for the time × diversity interaction). CWDC was

significantly lower in monocultures (least-squares mean of 0.010 kg C m^2 yr^{-1}) than in all mixtures, with

0.012, 0.024, and 0.021 kg C m^2 yr^{-1}, for 2-, 3- and 5-species mixtures, respectively. In contrast, ΔCRC

increment did not significantly differ between monocultures and mixtures (Table 2, Fig. 2).

Fig. 2 Mean changes in aboveground tree C (ΔAGC), coarse root C (ΔCRC) and coarse woody debris C (CWDC)
over time and with diversity. The analyses considered three time-intervals, early (p1), mid (p2), and late period (p3).
Coloured bars show means and error bars standard errors of the mean for the examined C stocks and fluxes calculated
as detailed in Eq. 1 and Fig. S1.

Table 2 Mixed-effects ANOVAs on changes in C stocks and fluxes over time. The analyses considered up to three
time-intervals, early, mid, and late period across a total of 16 years, see the timeline in Supplementary Fig. S1 for
details. Abbreviations are as in Figure 1. Entries in the table are F and p values, with p <0.05 *, p< 0.01 **, p<0.001
***, and ns = not significant. Significant effects printed in bold. For root decomposition and soil respiration, ANOVA
results are shown.

C stock/flux	Diversity	Time	Time × Diversity
ΔAGC	**5.07***	**11.25 *****	1.67 ns
ΔCRC	1.11 ns	2.74 ns	0.72 ns
CWDC	**3.30 ***	0.23 ns	2.01 ns
HerbaceousC	0.98 ns	**47.10 *****	0.37 ns
LitterC	0.81 ns	**100.74 *****	**6.25 *****
ΔSOC$_3$	0.92 ns	**29.10 *****	1.61 ns
ΔSOC$_4$	0.67 ns	**44.57 *****	**5.35 ****
ΔSOC	0.62 ns	0.01 ns	2.55 ns
Litter decomposition	0.71 ns	**30.32 *****	**5.15 ***
Root decomposition	0.69 ns	-	-
Microbial biomass	0.81 ns	**46.32 *****	0.90 ns
Soil respiration	0.45 ns	-	-
Canopy opening	2.06 ns	**31.73 *****	1.29 ns

We further compared the strongest aboveground C fluxes to the soil (herbaceousC and litterC, which were

an order of magnitude higher than CWDC; Fig. 1) with observed changes in ΔSOC (Fig. 3). Across the

plantation, herbaceousC did not vary with diversity but decreased significantly with time, with 0.201, 0.105,

and 0.125 kg C m^2 yr^{-1} for the early, mid, and late period, respectively (Table 2, Fig. 3). Conversely, litterC

increased significantly over time, with diversity effects depending on the period of plantation development

(Table 2): In the early period, litterC was lowest and similar across diversity levels, while in the mid and

late period litterC was lowest in monocultures with 0.149 and 0.141 kg C m^2 yr^{-1} and highest in 5-species

mixtures, with 0.225 and 0.232 kg C m^2 yr^{-1} (Table 2, Fig. 3).

The predominately tree-derived ΔSOC_3 showed a significant change over time but not with diversity (Table

2): in the mid period all plots – irrespective of their diversity – showed a net increase in ΔSOC_3 while most

plots did not show any SOC_3 loss or increment in the late period (Fig. 3). A notable exception are the 5-

species mixtures, which tended to lose SOC_3 in the late period (Fig. 3). The indication of enhanced SOC_3

loss in 5-species mixtures in the late period coincided with the slowest leaf litter decomposition in this

treatment (Fig. 4). Overall, leaf litter decomposition changed significantly over time and with diversity

(Table 2; Fig. 4). Litter decomposition was slowest in the early and fastest in the mid period but did not

respond to diversity in either period; in contrast, we observed a pronounced decline in leaf litter

decomposition with diversity in the late period, particularly in 5-species mixtures (note that decomposition

was only measured in 3- and 5-species mixtures in the mid and late period). A similar decrease in

decomposition with diversity – but non-significant – was observed for root decomposition (Table 2, Fig. 4;

note that root decomposition was only measured in the mid period).

ΔSOC_4, which is associated mainly with the C_4 grasses that existed before the plantation establishment,

varied significantly in response to both time and the time by diversity interaction (Table 2). Between 2001

and 2011 the reduction in C_4 in the upper 0–10 cm of soil was around twice as fast as between 2011 and

2017 with least-square means of -0.13 kg C m^2 yr^{-1} and -0.055 kg C m^2 yr^{-1}, respectively (Fig. 3). The

significant time by diversity interaction was driven by the three-species mixtures where the decrease in

ΔSOC_4 was strongest in the mid period and disappeared in the late period while the other treatments

experienced continued loss in ΔSOC_4 in the late period (Fig. 3). Overall, reductions in predominately

pasture-derived SOC_4 were larger than any observed gains in tree-derived SOC_3 resulting in a net negative

SOC-balance of reforestation (Fig. 3).

Fig. 3 Mean changes of dominant aboveground C fluxes to the soil with time and diversity and corresponding observed changes in soil organic carbon (ΔSOC). Coloured bars show means and error bars standard errors of the mean of herbaceous C and leaf litter C and changes in C_3 and C_4-derived SOC (SOC_3 and SOC_4). The analyses considered two time-intervals, mid (p2), and late period (p3), as SOC data was only measured in these two periods (Supplementary Fig. S1). The examined SOC changes were calculated as detailed in Eq. 1 and Fig. S1.

Fig. 4 Mean changes in leaf litter and root decomposition with time and diversity. Coloured points show means and error bars standard errors of the mean. The analyses considered three time-intervals, early (p1), mid (p2), and late period (p3). Leaf litter decomposition but not root decomposition responded significantly to tree diversity (Table 2).

Finally, none of the remaining variables, including microbial biomass, soil respiration and canopy opening responded significantly to diversity (Table 2), even though canopy opening tended to be lower in 5-species mixtures compared to monocultures (F = 2.06, p = 0.14), particularly in the mid- and late period. However, for soil microbial biomass and canopy opening, for which we had repeated measurements, we observed pronounced temporal changes (Table 2). Soil microbial biomass increased significantly from 315 to 453 µg C_{mic} g soil dw^{-1} from the mid to the late period. Not surprisingly, canopy opening, and thus light transmission, declined strongly with progressing stand development from a least-squares mean of 47.8% in the early to 21.1% in the mid period to than increase slightly again to 27.2% in the late period.

Effects of diversity on C stock-flux relationships

Using structural equation models, we explored the effect of tree diversity on linkages amongst C

stocks and fluxes above- and belowground (Fig. 5). In both periods of plantation development (mid and

late period), tree diversity significantly decreased canopy opening and increased ΔAGC with standardized

path coefficients of -0.41 and 0.42 in the mid period and -0.4 and 0.71 in the late period (Fig. 5). Tree

diversity effects on ΔAGC thus increased by ~70% from the mid to the late period. Canopy opening exerted

a significant positive effect on herbaceousC but decreased litterC with standardized path coefficient of 0.57

and -0.40 in the mid period and 0.52 and -0.43 in the late period. ΔAGC significantly increased both litterC

and CWDC with standardized path coefficients of 0.51 and 0.68 in the mid period and 0.53 and 0.62 in the

late period. Hence, we observed a consistent control of diversity on aboveground C stocks and fluxes and

their linkages across time. Aboveground, tree diversity led to a net increase of C fluxes to the soil through

its indirect effects on herbaceousC, litterC and CWDC of -0.23, 0.38, 0.29 and -0.21, 0.55, 0.44 respectively

for the mid- and late period (see Lefcheck, 2016 for calculation of indirect effects).

In contrast, we observed few significant linkages between aboveground and belowground C stocks

and fluxes and only one indirect effect of diversity on belowground changes in C stocks (Fig. 5). In the mid

and late period, the significant effect of litterC on belowground ΔC stock (expressed as sum of ΔCRC and

predominately tree-derived SOC fraction, $ΔSOC_3$) had a standardized path coefficient of 0.91 and 0.71,

respectively. The indirect effect of tree diversity on belowground ΔC stock via litterC was 0.36 and 0.39

for the mid and late period, respectively. SEMs expressing belowground ΔC stock using SOC instead of

SOC_3 yielded similar results, except for a disappearance of the significant effect of litterC in the late period

and overall lower explained variation (R^2) in ΔC stock (Fig. S5). Overall, belowground changes in C stocks

were largely disconnected from the diversity-controlled C stock and flux network observed aboveground.

Fig. 5 Direct and indirect effects of tree diversity on C stocks and fluxes. The SEMs partition potential tree diversity effects on C stocks and fluxes into effects mediated via canopy space filling (expressed as canopy opening) and via aboveground tree C changes (expressed as ΔAGC), which are expected to influence herbaceous C, leaf litter C and coarse woody debris C (CWDC). These latter C fluxes are, in turn, hypothesized to influence changes in belowground C, expressed as the summed change in coarse root C (ΔCRC) and predominately tree-derived soil organic C (ΔSOC$_3$). SEMs were fit for the mid period (panel A, P2) and the late period (panel B, P3) with all variables being calculated as detailed in Eq. 1 and Fig. S1. The SEMs fit the data well (Fisher's $C = 20.17$, df $= 20$, $P = 0.45$, $n = 22$ plots for (A); Fisher's $C = 21.17$, df $= 18$, $P = 0.27$, $n = 22$ plots for (B)). Examined variables are shown as boxes and relationships as directional arrows with significant positive effects in blue, significant negative effects in red, and nonsignificant effects in dashed gray. The hypothesis-driven conceptual model is shown in Fig. S3. For each significant relationship, standardized path coefficients are shown next to each path with path-width scaled according to coefficient size and asterisks indicating the significance level (*P < 0.05, **P < 0.01, and ***P < 0.001). Bidirectional grey arrows show significant partial correlations. The variation explained in each variable (R^2) is shown below the variable name. The green and brown font indicates above- and belowground variables.

Discussion

As the first tropical site of TreeDivNet, the Sardinilla experiment provides a unique opportunity to evaluate temporal changes in C stocks and fluxes in the neotropics. Consistent with our hypothesis (H1), we observed that tree diversity can increase C stocks and fluxes. Specifically, we noted a remarkable increase in AGC stocks driven by tree diversity, with an average 57% increase in AGC in five-species mixtures compared to monocultures after 16 years (35.7±1.8 *vs* 22.8±3.4 Mg C ha^{-1}; Fig. 1). This observed effect aligns with a recent meta-analysis demonstrating that mixture plots sequestered up to 70% more carbon in aboveground biomass than monoculture ones (Warner et al., 2023). Comparing our results with chronosequences of secondary forests in Panama and the wider neotropics, we found that our C stock

estimates after 16 years (35.8 and 22.6 Mg C ha^{-1} stored respectively in trees and SOC (0–10 cm)) are

comparable. For instance, Neumann-Cosel et al. (2011), working like us in the Panama Canal watershed,

reported aboveground ranges of 20.55 Mg C ha^{-1} to 56.45 Mg C ha^{-1} with SOC in the first 0-10 cm adding

27.5±3.1 Mg C ha^{-1}. Similarly, Gardon et al. (2020) estimated that actively restored secondary forests in

Brazil accumulated approximately 100 Mg biomass yr^{-1} after 16 years, roughly equivalent to 50 Mg C ha^{-1}

, and akin to the AGC of our most productive mixture plot. Overall, these findings contribute to our

understanding of the build-up of C stocks in tropical forest restoration and emphasize the importance of

considering tree diversity in such initiatives.

The long-term C balance of reforestation in the face of progressing climate change may depend more on

forest stability and thus C residence time within the forest than on average C accumulation rates. Here, we

understand stability as a forest's ability to maintain functioning over time despite repeated perturbations,

such as climate extremes (Schnabel et al., 2021), which is broadly consistent with the insurance hypothesis

(Yachi and Loreau, 1999). In the Sardinilla experiment, observed climate extremes left the ecosystem little

time to recover between perturbations. The extreme wet spell of 2010 was followed in 2015 by a severe

drought (Fig. S2; see also Browne et al., 2021; Detto et al., 2018), and a hurricane in 2016. As Bhaskar et

al. (2018) pointed out, it is important to understand forest resilience in the context of constant climatic

disturbances. While our study design does not allow us to disentangle the effects of climate extremes from

those of stand development on tree diversity effects, it does allow us to test whether positive tree diversity

effects persist or even strengthen over time despite repeated climate extremes, which triggered growth

reductions and elevated tree mortality (Hutchison et al., 2018; Schnabel et al., 2019). Consistent with our

hypothesis (H2), and despite these repeated extremes, we uncovered significant positive effects of tree

diversity on all the aboveground C stocks and fluxes (Fig. 5). Remarkably, the positive tree diversity effect

not only persisted but tended to strengthen over time, at least for ΔAGC. Our previous findings (Hutchison

et al., 2018; Schnabel et al., 2019) indicated that tree diversity increased the stability of tree productivity

by enhancing growth, buffering temporal variations in growth, and reducing mortality vis-à-vis

monocultures. Although two- and three-species mixtures were the most productive in the early period of

the Sardinilla experiment (e.g. Healy et al., 2008; Scherer-Lorenzen et al., 2007), our results demonstrate

that five-species mixtures outperformed the less diverse mixtures in later years, as already observed in

previous reports (Guillemot et al., 2020; Schnabel et al., 2019). Here, we moved beyond these earlier studies

and similar findings in other ecosystems (e.g. Jucker et al., 2014; Schnabel et al., 2021), which focussed on

single ecosystem functions, and leveraged a unique dataset on multiple C stocks and fluxes and their

interrelationships. This approach allowed us to adopt an integrated ecosystem perspective on C stability in

mixed compared to monospecific planted forests. C stability is particularly important considering that tree

diversity increased C stocks and fluxes only aboveground, where C is particularly susceptible to climate-

driven forest disturbances.

Using structural equation modelling we showed that the tree diversity directly or indirectly effected all the

aboveground C stocks and fluxes that we measured (Fig. 5), suggesting that the aboveground components

of the C cycle are linked. Canopy opening, a proxy for canopy space filling, played a central role in our

SEMs with denser foliage at high diversity enhancing litterC but reducing herbaceousC. Tree architecture

apparently plays a crucial role in explaining these effects of tree diversity on the aboveground system. After

16 years of growth, trees growing in mixture allocated a higher proportion of their biomass to branches

(Guillemot et al., 2020), which is consistent with higher canopy space filling reported in other tree diversity

experiments (Kunz et al., 2019; Williams et al., 2017). Similarly, tree productivity (captured here as ∆AGC)

increased with tree diversity, as has been previously reported from our (Guillemot et al., 2020; Schnabel et

al., 2019) and other tree diversity experiments (Guerrero-Ramírez et al., 2017). Enhanced tree productivity

at high diversity, in turn, enhanced both litterC and CWDC and thus C fluxes to the soil. Likely drivers of

these observed positive diversity effects aboveground are complementary species interactions in mixtures,

such as higher community-level light capture or complementary water and nutrient uptake from different

soil layers reported in Sardinilla (Oelmann et al., 2010; Sapijanskas et al., 2014; Schwendenmann et al.,

2015; Zeugin et al., 2010). Litter manipulation experiments support the idea of significant leaf litter-driven

carbon and nutrient fluxes to the soil. Long-term experiments in Barro Colorado Islands Nature Monument

in Panama removed litter from forest plots to add it to other identical plots, allowing to compare plots with

litter removal and litter addition (Cusack et al., 2018; Sayer et al., 2024). After ten and 17 years of

manipulation, soil C was significantly higher in the 0–5 cm layer in the litter addition plots (Cusack et al.,

2018), while tree growth significantly declined in litter removal plots (Sayer et al., 2024). The authors

explained their results by the fast carbon cycling and high C input into the soil in tropical forests. Our SEMs

confirmed the positive link between litterC and the predominately tree-derived fraction of SOC (SOC_3) but

no further linkages of SOC_3 with tree diversity nor herbaceousC or CWDC (Fig. 5). This means that the

surplus of C fluxes to the soil in the high-diversity mixtures was largely not incorporated into the mineral

soil phase, and thus, the multiple direct and indirect relationships of aboveground C stocks and fluxes

largely did not extend belowground.

Overall, the Sardinilla planted forest gained significant C aboveground, but we observed a loss of SOC in

the 0–10 cm layer associated with a reduction in both bulk density and SOC concentration (Table S2). This

loss occurred regardless of tree species richness levels, indicating that tree diversity may only sometimes

exert a significant effect on SOC, as noted also in another temperate tree diversity experiment (Martin-

Guay et al., 2022). Because the planted forest was established on an active pasture (Scherer-Lorenzen et

al., 2005), we suggest that the removal of cows likely minimized compaction which resulted in decreased

bulk density (Blanco Sepúlveda and Nieuwenhuyse, 2011). Several studies have identified the

misrepresentation of changes in SOC mass associated with sampling to a fixed depth where there have been

significant changes in bulk density to that depth (e.g. von Haden et al., 2020). Although variable among the

treatments and the two sampling periods, comparison of relative changes in SOC concentration and bulk

density suggest that about half of the apparent loss in SOC mass is associated with a loss of SOC

concentration and the other half with a decrease in bulk density (Table S2) and thus the loss of SOC mass

in the plantation may be an overestimate (Fig. 3). However, SOC concentrations in the 10 – 50 cm depth in

the pasture prior to the establishment of the Sardinilla experiment ranged from 1 to 2% and a $\delta^{13}C$ value

between -17 and -21‰, suggesting a strong proportion of SOC_4, and thus there may be further losses in

SOC from the subsoil (Moore et al., 2018), as also suggested by (Quartucci et al., 2023). The literature fails

to reach a consensus about how reforestation affects SOC (Laganière et al., 2010), with climatic zone,

species planted, clay content, past land use, and soil parent material all affecting the SOC balance of forest

regrowth (Araujo et al., 2017; Wallwork et al., 2022). For example, in Brazil's Atlantic forest, SOC stock

(0–10 cm) of 5-year old re-growing forests was half that of the remnant forest and similar to that of pastures

(Zanini et al., 2021) while another Brazilian study (Camargo et al., 1999) reported similar SOC stocks at

0–10 cm for forest, secondary forest fallows, and pasture even though AGC varied 44-fold. Marín-Spiotta

et al. (2009), who reported no changes in SOC during secondary forest establishment in Puerto Rico,

explained this by a loss of pasture-derived SOC_4 which counterbalanced tree-derived gains in SOC_3.

Finally, in a nearby Panamanian site, SOC stocks did not vary along a chronosequence of secondary forests

(Neumann-Cosel et al., 2011). We propose that Sardinilla's clay-rich Cambisols, Tropudalfs and Vertisols

(average clay content reaches 65% (Moore et al., 2018)) might have amplified compaction related to grazing

while the establishment of trees likely loosened up the soil (Table S2) with follow-up effects like an increase

in mineralization and thus, increased C loss. Important gaps in the understanding of SOC cycling and

storage have restricted the ability to manage ecosystems to enhance soil C stocks (Wallwork et al., 2022).

In the case of Sardinilla, we suggest that the land-use change effect (partly) overruled the effects of tree

diversity and its C_3-derived C inputs to SOC stock. Given that high soil clay content is common in the

tropics (http://hydro.iis.u-tokyo.ac.jp/~sujan/research/gswp3/soil-texture-map.html; Rasmussen et al.,

2018) including soil C pool in the C assessments of forest restoration is likely to improve overall C storage

estimates (Quartucci et al., 2023).

The importance of forest restoration for ecosystem C storage, a key ecosystem service for climate

regulation, has not only been discussed in the scientific literature but has also triggered international

commitment to reforestation. As we work on understanding how forest C stocks are being rebuilt through

time at the ecosystem level, it is important to remain realistic about the potential of forest restoration to

contribute towards mitigating climate change. Sequestering C is a slow process (Baldocchi and Penuelas,

2019): the average yearly net CO_2 uptake in Sardinilla was 5.67 Mg CO_{2eq} ha^{-1} yr^{-1} or 1.54 Mg C ha^{-1} yr^{-1}.

To illustrate, we estimated the emissions from a single one-way flight between Frankfurt and Panama City

to 62.7 Mg CO_{2eq} using the ICAO carbon calculator. Hence, this flight demands a flux equivalent to that

sequestered by ~11 ha in the Sardinilla planted forest in one year. Some have suggested that the enthusiasm

for nature-based solutions risks putting excessive pressure on land use. For example, Dooley et al. (2022)

estimated that countries' climate pledges for land-based carbon dioxide removal would demand 1.2 billion

ha of land, an area globally equal to that used to grow food. Our thorough ecosystem-level analysis of C

stocks and fluxes sheds some light on the challenges of using active reforestation projects to compensate

for emissions. We note that Griscom et al. (2017) used higher sequestration potentials of 2.8 to 4.7 Mg C

ha^{-1} yr^{-1} to estimate the potential of reforestation to act as a nature-based solution for climate mitigation.

Therefore, while nature-based solutions are undeniably important for C sequestration and other ecosystem

benefits, such as biodiversity and water regulation (Griscom et al., 2017; Seddon, 2022), co-benefits and

potential trade-offs should be carefully assessed (Schuldt et al., 2023). Mixed planted forests as a nature-

based solution may not only enhance carbon stocks and fluxes vis-à-vis monocultures, as we show here,

but also decrease the restored forests susceptibility to stress and disturbances and, thereby, increase C

permanence (Anderegg et al., 2020) while also providing higher levels of biodiversity and a broader range

of ecosystem services (Messier et al., 2021).

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686 Author contributions

687 CP and FS conceived the goals and aims of the study and coordinate the Sardinilla experiment. FS, CP, JG,

688 KEB, SC, NE, and CW developed the methodology. CP, FS, JG, MB, SC, NE, TG, NRGR, ITH, CM, LM,

689 JM, TM, YO, MSL, LS, and AW collected and curated the data. FS and CP analysed the data. FS created

690 the figures. FS and CP wrote the original manuscript draft and all authors contributed to reviewing and

691 editing the manuscript.

693 Conflict of interest statement

694 The authors declare no conflict of interest.

696 Data availability

The data that support the findings of this study are openly available in Dryad at http://doi.org/tbd. Data will

be made openly available upon acceptance.

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