

1 **Title:** The fate of carbon in a mature forest under carbon dioxide enrichment

2 M. Jiang¹, B.E. Medlyn¹, J.E. Drake^{1,2}, R.A. Duursma¹, I.C. Anderson¹, C.V.M. Barton¹,
3 M.M. Boer¹, Y. Carrillo¹, L.Castañeda-Gómez¹, L. Collins^{1,3,4}, K.Y. Crous¹, M.G. De
4 Kauwe^{5,6,7}, B.M. dos Santos^{8,9}, K.M. Emmerson¹⁰, S.L. Facey¹, A.N. Gherlenda¹, T.E.
5 Gimeno^{1,11,12}, S. Hasegawa^{1,13}, S.N. Johnson¹, C.A. Macdonald¹, K. Mahmud^{1,14}, A.
6 Kännaste¹⁵, B.D. Moore¹, L. Nazaries¹, E.H.J. Neilson^{8,9}, U.N. Nielsen¹, Ü. Niinemets¹⁵, N.J.
7 Noh^{1,16}, R. Ochoa-Hueso^{1,17}, V.S. Pathare^{1,18}, E. Pendall¹, J. Pihlblad¹, J. Pineiro^{1,19}, J.R.
8 Powell¹, S.A. Power¹, P.B. Reich^{1,20}, A.A. Renchon¹, M. Riegler¹, R. Rinnan²¹, P. Rymer¹,
9 R.L. Salomón²², B.K. Singh^{1,23}, B. Smith^{1,24}, M.G. Tjoelker¹, J.K.M. Walker¹, A. Wujeska-
10 Klaue¹, J. Yang¹, S. Zaehle²⁵, and D.S. Ellsworth¹

11
12 **Affiliation:**

13 ¹Hawkesbury Institute for the Environment, Western Sydney University, Locked Bag 1797,
14 Penrith, NSW, 2751, Australia

15 ²Department of Sustainable Resources Management, College of Environmental Science and
16 Forestry, State University of New York, Syracuse, NY 13210, USA

17 ³Department of Ecology, Environment and Evolution, La Trobe University, Bundoora, VIC
18 3086, Australia

19 ⁴Arthur Rylah Institute for Environmental Research, Department of Environment, Land,
20 Water and Planning, PO Box 137, Heidelberg, VIC 3084, Australia

21 ⁵ARC Centre of Excellence for Climate Extremes, University of New South Wales, Sydney,
22 NSW 2052, Australia

23 ⁶Climate Change Research Centre, University of New South Wales, Sydney, NSW, 2052,
24 Australia

25 ⁷Evolution and Ecology Research Centre, University of New South Wales, Sydney, NSW,
26 2052, Australia

27 ⁸Plant Biochemistry Laboratory, Department of Plant and Environmental Sciences, University
28 of Copenhagen, Thorvaldsensvej 40, 1871 Frederiksberg C, Copenhagen, Denmark

29 ⁹VILLUM Research Center for Plant Plasticity, University of Copenhagen, Thorvaldsensvej
30 40, 1871 Frederiksberg C, Copenhagen, Denmark

31 ¹⁰Climate Science Centre, CSIRO Oceans & Atmosphere, Aspendale, VIC 3195, Australia

32 ¹¹Basque Centre for Climate Change, Leioa, 48940, Spain

33 ¹²Ikerbasque, Basque Foundation for Science, 48008 Bilbao, Spain

34 ¹³Department of Forest Ecology and Management, Swedish University of Agricultural
35 Sciences (SLU), Umeå, SE-90183, Sweden

36 ¹⁴Department of Geography, Indiana University, Bloomington, IN 47405, USA

37 ¹⁵Estonian University of Life Sciences, Kreutzwaldi 1, 51006, Tartu, Estonia

38 ¹⁶Forest Technology and Management Research Center, National Institute of Forest Science,
39 Gyeonggi-do, 1186, Korea

40 ¹⁷Department of Biology, IVAGRO, University of Cádiz, Campus de Excelencia
41 Internacional Agroalimentario (CeIA3), Campus del Río San Pedro, 11510 Puerto Real,
42 Cádiz, Spain

43 ¹⁸School of Biological Sciences, Post Office Box 646340, Washington State University,
44 Pullman, WA 99164-6340, USA

45 ¹⁹Division of Plant and Soil Sciences, West Virginia University, Morgantown, WV, USA

46 ²⁰Department of Forest Resources, University of Minnesota, St Paul, Minnesota, 55108, USA

47 ²¹Terrestrial Ecology Section, Department of Biology, University of Copenhagen,
48 Universitetsparken 15, DK-2100, Copenhagen, Denmark

49 ²²Laboratory of Plant Ecology, Faculty of Bioscience Engineering, Ghent University,
50 Coupure links 653, 9000 Ghent, Belgium

51 ²³Global Centre for Land Based Innovation, Western Sydney University, Building L9,
52 Locked Bag 1797, Penrith South, NSW, 2751, Australia

53 ²⁴Department of Physical Geography and Ecosystem Science, Lund University, 22362, Lund,
54 Sweden

55 ²⁵Max Planck Institute for Biogeochemistry, Hans-Knöll-Str. 10, 07745 Jena, Germany

56 **Abstract**

57 Atmospheric carbon dioxide enrichment (eCO₂) can enhance plant carbon uptake and
58 growth^{1,2,3,4,5}, thereby providing an important negative feedback to climate change by slowing
59 the rate of increase of the atmospheric CO₂ concentration⁶. While evidence gathered from
60 young aggrading forests has generally indicated a strong CO₂ fertilization effect on biomass
61 growth^{3,4,5}, it is unclear whether mature forests respond to eCO₂ in a similar way. In mature
62 trees and forest stands^{7,8,9,10}, photosynthetic uptake has been found to increase under eCO₂
63 without any apparent accompanying growth response, leaving an open question about the fate
64 of additional carbon fixed under eCO₂^{4,5,7,8,9,10,11}. Here, using data from the first ecosystem-
65 scale Free-Air CO₂ Enrichment (FACE) experiment in a mature forest, we constructed a
66 comprehensive ecosystem carbon budget to track the fate of carbon as the forest responds to
67 four years of eCO₂ exposure. We show that, although the eCO₂ treatment of ambient +150
68 ppm (+38%) induced a 12% (+247 g C m⁻² yr⁻¹) increase in carbon uptake through gross
69 primary production, this additional carbon uptake did not lead to increased carbon
70 sequestration at the ecosystem level. Instead, the majority of the extra carbon was emitted
71 back into the atmosphere via several respiratory fluxes, with increased soil respiration alone
72 accounting for ~50% of the total uptake surplus. Our results call into question the
73 predominant thinking that the capacity of forests to act as carbon sinks will be generally
74 enhanced under eCO₂, and challenge the efficacy of climate mitigation strategies that rely on
75 ubiquitous CO₂ fertilization as a driver of increased carbon sinks in global forests.

76

77 **Main text**

78 Globally, forests act as a large carbon sink, absorbing a significant portion of the
79 anthropogenic CO₂ emissions^{1,12}, an ecosystem service that has tremendous social and

80 economic value. Whether mature forests will remain carbon sinks into the future is of critical
81 importance for aspirations to limit climate warming to no more than 1.5 °C above pre-
82 industrial levels¹³. Free-Air CO₂ Enrichment (FACE) experiments provide an opportunity to
83 determine the capacity of ecosystems to sequester carbon under the higher atmospheric CO₂
84 concentrations expected in the future^{3,4,5,7,8,10,11}. Evidence gathered from the four first-
85 generation forest FACE experiments, which all measured responses of rapidly-growing
86 young forest plantations, has generally indicated a strong CO₂ fertilization effect on biomass
87 growth^{3,4}. This CO₂ fertilization effect has been hypothesized to be one of the largest drivers
88 of the terrestrial carbon sink and its acceleration in recent decades¹⁴, potentially accounting
89 for up to 60% of present-day terrestrial carbon sequestration². However, younger trees are
90 generally more responsive to rising CO₂ than mature trees¹¹, potentially because nutrient
91 limitation increases with stand age¹⁵. Thus, extrapolating evidence collected from these
92 experiments may be argued to provide an upper limit on how much carbon can be stored by
93 global forests under eCO₂¹⁶. Evidence from experiments with older trees on nutrient-poor
94 soils suggests that although eCO₂ increases leaf photosynthesis to a similar degree as in
95 young forests, stimulation of biomass growth and carbon storage may be lower or
96 absent^{7,8,9,10}. Reconciling these conflicting observations is a crucial step towards quantifying
97 the carbon sequestration capacity of mature forests in the future. It requires that we identify
98 the fate of the extra carbon fixed under eCO₂ in mature forests, which are expected to be
99 closer to a state of equilibrium between carbon uptake and turnover, compared to young
100 aggrading stands.

101

102 The *Eucalyptus* FACE (EucFACE) experiment is the world's first replicated, ecosystem-scale
103 mature forest FACE experiment (Extended Data Figure 1, 2). It is located in a warm-
104 temperate evergreen forest that has remained undisturbed for the past 90 years, is dominated

by the regionally widespread tree *Eucalyptus tereticornis* and has an understorey composed principally of native grasses and shrubs. The low-fertility soil has been shown to limit tree growth in an adjacent phosphorus-fertilization experiment¹⁷. Seven ecosystem-scale models were used to predict the eCO₂ response at EucFACE in advance of the experiment¹⁸, highlighting three alternative hypotheses for the expected ecosystem response based on plausible assumptions incorporated in different models¹⁹. These hypotheses were: (i) enhanced photosynthesis under eCO₂ would lead to increased biomass accumulation; (ii) eCO₂-induced increase in photosynthesis would be directly down-regulated by limited nutrient availability; or (iii) eCO₂-induced increase in photosynthesis would lead to increased autotrophic respiration¹⁸. This range of predictions among a suite of well-tested models indicated a prognostic knowledge gap as to how the carbon cycling of mature forests would respond to the expected rise in atmospheric CO₂ concentration¹¹, which is crucial to resolve in the face of future carbon-climate uncertainty²⁰.

To date, both canopy trees and understorey plants at EucFACE have shown increased rates of leaf photosynthesis but the canopy trees showed no significant increase in aboveground biomass growth under eCO₂⁷, reflecting a similar lack of response observed in other eCO₂ experiments on mature trees^{8,9,10}. Incorporating leaf-scale gas exchange measurements into a process-based tree stand model, it was estimated that the observed +19% stimulation of light-saturated overstorey leaf photosynthesis⁷ corresponded to a +11% stimulation of whole-canopy gross primary production (GPP) in response to eCO₂²¹. However, the probable fate of the extra carbon fixed under eCO₂ remained undetermined. Where did the extra carbon go?

To answer this question, we compiled measurements on all major carbon pools and fluxes collected over four years of experimental treatment (2013-2016), including individual and

aggregated biomass and associated fluxes measured or inferred from plants, litter, soil, microbes, and insects, and constructed an ecosystem carbon budget (Figure 1) under both ambient (aCO₂) and eCO₂ conditions (+150 ppm). We first confirmed mass balance of the ecosystem carbon budget by checking agreement between independent estimates of GPP and soil respiration (R_{soil}) derived from separate data streams (Extended Data Figure 3; see Methods). For GPP of the aCO₂ plots, we confirmed that a process-based model estimate of overstorey and understorey GPP ($2059 \pm 211 \text{ g C m}^{-2} \text{ yr}^{-1}$), driven by site-specific meteorology and treatment-specific physiological data, broadly agreed with the sum of data-driven estimates of net primary production (NPP) and autotrophic respiration ($2068 \pm 61 \text{ g C m}^{-2} \text{ yr}^{-1}$). The carbon-use efficiency (NPP/GPP) of this mature forest was estimated to be 0.31 ± 0.03 , which is on the low end of global forest estimates, but consistent with studies that have observed this ratio to decline with stand age²² (Extended Data Figure 2). We further confirmed carbon mass balance for R_{soil} of the aCO₂ plots by comparing soil chamber-based estimates ($1097 \pm 86 \text{ g C m}^{-2} \text{ yr}^{-1}$) with the sum of litterfall and independently estimated root respiration ($1086 \pm 14 \text{ g C m}^{-2} \text{ yr}^{-1}$), assuming no change in soil carbon pool (see Methods). This agreement between independent estimates of components of the ecosystem carbon budget gives confidence that our measurements captured the pools and fluxes of carbon with low aggregate uncertainty and hence allow us to infer the fate of the extra carbon fixed under eCO₂.

To accommodate the inherent pre-treatment plot differences (see Methods), we normalized the CO₂ responses across plots by using a linear mixed-model with plot-specific pre-treatment leaf area index as a covariate^{23,24}. The non-normalized eCO₂ responses are provided in Extended Data Figure 4, and generally confirm the findings but with larger uncertainty. Our normalized responses (Figure 2, Extended Data Figure 5) showed that eCO₂ induced an

average of 12% increase ($+247 \pm 195 \text{ g C m}^{-2} \text{ yr}^{-1}$, mean \pm one standard deviation) in carbon uptake, including contributions of overstorey ($+192 \pm 193 \text{ g C m}^{-2} \text{ yr}^{-1}$) and understorey GPP ($+55 \pm 21 \text{ g C m}^{-2} \text{ yr}^{-1}$). The fate of this additional carbon entering the system under eCO_2 was primarily traced to an increase in R_{soil} ($+128.8 \pm 116.7 \text{ g C m}^{-2} \text{ yr}^{-1}$, or 52% of the carbon uptake surplus), followed by a smaller increase in tree stem respiration (R_{stem} ; $+40.0 \pm 43.6 \text{ g C m}^{-2} \text{ yr}^{-1}$, or 16% of the carbon uptake surplus). In comparison, the increase in total NPP ($+67.3 \pm 12.7 \text{ g C m}^{-2} \text{ yr}^{-1}$, or 28% of the carbon uptake surplus) corresponded to a smaller increase in storage of the total carbon pools at the ecosystem-level (ΔC_{pools} ; $+31.6 \pm 188.8 \text{ g C m}^{-2} \text{ yr}^{-1}$, or 12.8% of the carbon uptake surplus, Extended Data Figure 6). There was thus little evidence of additional carbon accumulation under eCO_2 in this mature forest ecosystem. We then compared three alternative methods (see Methods) of estimating net ecosystem production (NEP; Figure 3). All three indicated that the ecosystem remained close to carbon-neutral under ambient CO_2 over the experimental period (mean \pm SD for the methods: 28 ± 225 , 21 ± 129 , $-73 \pm 50 \text{ g C m}^{-2} \text{ yr}^{-1}$, respectively), and that eCO_2 of +150 ppm did not result in statistically significant increases in ecosystem carbon storage (109 ± 258 , -19 ± 171 , $-42 \pm 262 \text{ g C m}^{-2} \text{ yr}^{-1}$, respectively). However, the variability reported here means that we cannot fully rule out the possibility of additional carbon storage under eCO_2 , but we stress that our individual and aggregated responses consistently suggest a lack of CO_2 response in this mature forest (Figure 2 & 3, Extended Data Figure 5).

The relatively small but positive NPP response to eCO_2 was mainly driven by the understorey aboveground NPP response (NPP_{ua} ; $+50.3 \pm 17.9 \text{ g C m}^{-2} \text{ yr}^{-1}$), which was 75% of the net NPP response (Figure 2). However, this significant NPP_{ua} response did not result in an equivalent eCO_2 effect on understorey aboveground biomass increment ($+27.2 \pm 29.7 \text{ g C m}^{-2} \text{ yr}^{-1}$), suggesting a possible higher understorey biomass turnover under eCO_2 . Smaller fluxes,

often neglected in other ecosystem carbon budgets, such as leaf consumption by insect herbivores (NPP_{ins} ; 25.5 ± 4.3 vs. $27.8 \pm 6.3 \text{ g C m}^{-2} \text{ yr}^{-1}$, aCO_2 vs. eCO_2 mean \pm SD), insect frass production (Frass; 10.5 ± 1.8 vs. $11.4 \pm 2.6 \text{ g C m}^{-2} \text{ yr}^{-1}$), vegetation volatile carbon emission (VC; 2.63 ± 0.18 vs. $2.45 \pm 0.13 \text{ g C m}^{-2} \text{ yr}^{-1}$), net ecosystem methane uptake (CH_4 ; 0.18 ± 0.0009 vs. $0.19 \pm 0.0003 \text{ g C m}^{-2} \text{ yr}^{-1}$), and leaching of dissolved organic carbon (DOC; 0.16 ± 0.017 vs. $0.17 \pm 0.024 \text{ g C m}^{-2} \text{ yr}^{-1}$), contributed to the closure of the overall ecosystem carbon budget (Figure 1; Extended Data Figure 3), but were not quantitatively important in explaining pathways of the carbon uptake surplus under eCO_2 (Figure 2, Extended Data Figure 5, Extended Data Figure 6).

Here we provide some of the first replicated experimental evidence on the probable fate of carbon under eCO_2 in intact mature forest. We found that increased R_{soil} accounted for ~50% of the extra photosynthate produced by plants under eCO_2 . It has been suggested that the increase in R_{soil} at EucFACE was likely a consequence of increased root and rhizosphere respiration^{25,26}, in contrast to other FACE sites where increased R_{soil} was attributed to enhanced soil organic matter decomposition (e.g. DukeFACE²⁷). Here, the eCO_2 -induced increase in R_{soil} was not accompanied by substantial changes in root respiration ($18.6 \pm 20.1 \text{ g C m}^{-2} \text{ yr}^{-1}$) or in carbon pools associated with fine roots ($+7.0 \pm 12.5 \text{ g C m}^{-2} \text{ yr}^{-1}$), microbes ($+1.9 \pm 3.5 \text{ g C m}^{-2} \text{ yr}^{-1}$), mycorrhizae ($+0.4 \pm 0.5 \text{ g C m}^{-2} \text{ yr}^{-1}$), leaf litter ($+27.1 \pm 38.6 \text{ g C m}^{-2} \text{ yr}^{-1}$) or soil ($-23.8 \pm 159.6 \text{ g C m}^{-2} \text{ yr}^{-1}$), suggesting that the additional carbon fixed under eCO_2 may have led to an enhanced carbon transport belowground and a rapid belowground turnover of this flux. Assimilation of these data into a carbon balance model supports this inference (Extended Data Figure 7, see Methods for details). An initial enhancement in nitrogen and phosphorus mineralization was observed²⁸, which suggested that the increased R_{soil} with eCO_2 could reflect soil organic matter priming with the potential to alleviate plant

nutrient stress in this low-phosphorus soil^{28,29}. However, the enhanced soil mineralization rate and associated increase in nutrient availability did not persist over time²⁸, indicating that this increased belowground carbon allocation and the rapid turnover of this flux was not effective in increasing phosphorus availability to the plants³⁰.

The ecosystem carbon budget presented here provides an opportunity to confront the three alternative hypotheses of the response of this system to eCO₂ treatment that emerged from model predictions made in advance of the experiment¹⁸. Our data do not support any of the three hypotheses. The eCO₂-induced increase in photosynthesis was not strongly down-regulated by low nutrient availability^{7,21}; nor did the eCO₂-induced additional carbon uptake lead to additional biomass accumulation, or enhanced aboveground respiration. These predictions reflect common mechanisms by which terrestrial vegetation models implement nutrient limitation of the eCO₂ response^{18,19,31,32}. In contrast, our results suggest a direct connection between plant photosynthesis and belowground activity (Extended Data Figure 7), in which increased belowground carbon allocation increased soil respiration at a rate that accounted for half of the extra carbon fixed under eCO₂ (Figure 2). Predictions made in advance of the experiment did not capture this additional belowground carbon flux, despite their general agreement with data on turnover rates of major carbon pools (Extended Data Figure 8). This increased soil respiration has been demonstrated by some models to be an important and often overlooked mechanism that reduces global soil carbon sequestration relative to estimates by many current models³³. As a consequence of including this rapid turnover of the increased belowground carbon allocation in terrestrial biosphere models, the time-lag in emitting some of the extra carbon via biomass accumulation and litterfall input into the soils may be reduced, thereby leading to faster cycling of carbon³⁴ and therefore possible different trajectories of carbon-climate predictions for the future.

230

231 A major form of land-based climate mitigation actions envisaged in the 2015 Paris
232 Agreement is to enhance forest biomass carbon stocks globally through the protection of
233 existing, largely mature, forests, and through afforestation of new areas. The mitigation
234 potential of forests lies in the accumulated stock of ecosystem carbon, not in the short-term
235 rate of forest photosynthesis. The probable fate of additional carbon determined in our study
236 (Figure 2) challenges the current thinking that all non-aggrading mature forests will
237 contribute to enhanced carbon sinks due to CO₂ fertilization³⁵, which further questions the
238 allowable CO₂ emission targets sourced from existing carbon cycle models^{13,36}. Given that
239 the effect of CO₂ fertilization may be one of diminishing returns over time¹⁴, the statistically
240 non-significant eCO₂ effect on NEP (Figure 3), if representative of nutrient-limited mature
241 forest ecosystems generally, suggests an even weaker carbon sink in the future, especially in
242 low-phosphorus systems such as EucFACE. Future research efforts should target a deeper
243 understanding of the nutrient-carbon feedbacks that likely constrain the carbon sink potential
244 of mature forests under eCO₂, and evaluate the implications of a potentially weaker terrestrial
245 land carbon sink in the development of robust mitigation strategies in the face of climate
246 change. More importantly, whilst the terrestrial carbon sink is integral to current strategies for
247 climate change mitigation, our results call for more active reductions of anthropogenic
248 emissions to meet the targets of the Paris Agreement.

Methods

EucFACE site description

The EucFACE facility (Extended Data Figure 1) is located in a mature evergreen *Eucalyptus* forest on an alluvial spodosol in western Sydney, Australia (33°36'S, 150°44'E). The site has been a remnant patch of native Cumberland Plain woodland since the 1880's and has remained unmanaged for at least the past 90 years, with *Eucalyptus tereticornis* Sm. as the dominant tree species (98% of the overstorey basal area). *Eucalyptus* trees occur naturally across Australia, accounting for 78% of native forest area in Australia³⁷ and are planted widely around the globe³⁸. Infrastructure for six large circular plots (490 m² each) was established in 2010. Starting on 18th September 2012, three plots were subjected to free-air CO₂ enrichment treatment using a computer-controlled pre-dilution method. The CO₂ concentrations at EucFACE were ramped up over a six-month period, increasing by +30 ppm every five weeks in discrete steps (+30, 60, 90, 120, and 150 ppm). The full elevated CO₂ treatment of +150 ppm started on 6th February 2013 during daylight hours over all days of the year. The site is characterized by a humid temperate-subtropical transitional climate with a mean annual temperature of 17.5°C and a mean annual precipitation of 800 mm (Figure S1). The soil is a Holocene alluvial soil of low fertility with low phosphorus content^{7,17}. Soil texture is a loamy sand (> 75% sand content) up to 50 cm in depth. From ca. 50 to 300 cm depth, soils are sandy clay loam, with > 30% silt and clay. Average bulk density is 1.39, 1.69 and 1.71 g cm⁻³ for depths of 0-10, 10-20 and 20-30 cm, respectively (Figure S2). Permanent groundwater depth is ~11 m below the soil surface³⁹. Understorey vegetation is a diverse mixture of 86 species including forbs, graminoids and shrubs⁴⁰. The dominant understorey species is *Microlaena stipoides*, a C₃ perennial grass that accounted for ~70% of herbaceous biomass and responded rapidly to rainfall variability⁴¹.

273

274 **Estimates of carbon pools and fluxes**

275 We estimated plot-specific carbon pools and fluxes at EucFACE over 2013-2016 (Extended
276 Data Table 1). We defined pools as a carbon reservoir and annual increments as the annual
277 changes in the size of each reservoir. We compartmentalized the ecosystem into 11 carbon
278 pools, namely overstorey leaf (C_{ol}), stem (C_{stem}), fine root (C_{froot}), coarse root (C_{croot}),
279 intermediate root (C_{iroot}), understorey aboveground (C_{ua}), soil (C_{soil}), microbe (C_{micr}),
280 mycorrhizae (C_{myco}), leaf litter (C_{lit}), and aboveground insect (C_{ins}) carbon pools, and reported
281 pool size in the unit of $g\ C\ m^{-2}$. We defined fluxes as components of the carbon flow through
282 the system, and report them in the unit of $g\ C\ m^{-2}\ yr^{-1}$. All annual incremental changes in
283 carbon pools were reported in $g\ C\ m^{-2}\ yr^{-1}$ with a symbol Δ . We converted estimates of
284 biomass into carbon content using variable-specific carbon fractions (f) defined in Extended
285 Data Table 2. Below we describe how each pool and flux was estimated.

286

287 Pools

288 **Soil carbon pool** (C_{soil} ; Figure S2) was estimated based on quarterly sampled soil carbon
289 content (oven-dried at 40 °C for 48 hours) and plot-specific soil bulk density at three depths
290 (0 - 10 cm, 10 - 20 cm, 20 - 30 cm). Out of the 15 dates when samples were taken, soil carbon
291 content below the top 10 cm of soil was measured on three dates. To obtain a more accurate
292 estimate of annual incremental change in soil carbon pool, we therefore reported soil carbon
293 pool for the top 10 cm only. There were no temporal and eCO_2 trends in soil carbon content
294 at deeper depths.

295

Overstorey leaf carbon pool (C_{ol} ; Figure S3) was estimated based on continuous measures of leaf area index (LAI) and specific leaf area (SLA, m^2 leaf area g^{-1} leaf DM), following $C_{ol} = LAI \times SLA \times f_{ol}$, where f_{ol} is a carbon fraction constant for overstorey leaves (Extended Data Table 2). Daily averages of plot-specific LAI were estimated based on the attenuation of diffuse radiation in a homogenous canopy²⁴. The number of observations varies between days, depending on the number of 30-minute cloudy periods. SLA was estimated based on time-series measures of leaf mass per area (LMA), and was then linearly interpolated to plot-specific daily values over time.

Stem carbon pool (C_{stem} ; Figure S4) was estimated based on tree-specific height and diameter at breast height (DBH) measurements, and an allometric scaling relationship derived for *E. tereticornis*^{7,42}. DBH changes were measured repeatedly at roughly monthly intervals, at 1.3 m height. Bark was periodically removed from under the dendrometer bands - this effect on DBH was considered by calculating biomass once per year using December data only. Stem biomass data were summed for each plot and averaged over the plot area to obtain ground-based estimates, and was then converted into C_{stem} using treatment-specific carbon fraction (Extended Data Table 2).

Understorey aboveground carbon pool (C_{ua} ; Figure S5) was estimated at 1-3 month intervals between February 2015 and December 2016 using non-destructive measurements of plant height obtained from stereo-photography⁴³. In each of the four $2m \times 2m$ understorey monitoring subplots within each plot, stereo photographs were collected using a Bumblebee XB3 stereo camera (Point Grey Research) mounted ~ 2.4 m above the ground surface and facing vertically downwards towards the center of the subplot. Stereo images were taken at dusk under diffuse light conditions to avoid measurement errors related to shadows from

trees and EucFACE infrastructure. On each sampling date, three sets of stereo photographs were taken in each subplot to produce a large number (i.e. 100,000 s) of understorey plant height estimates from which mean plant height (H_{mean} , in m) was calculated for each plot. Understorey aboveground biomass (B_{ua} , in kg m^{-2}) for each plot was predicted from H_{mean} using an empirical model developed for the grassy understorey vegetation at EucFACE ($B_{\text{ua}} = 1.72 \times H_{\text{mean}} - 0.05$)⁴³. The four subplot-level estimates were averaged to obtain a plot-level estimate of B_{ua} , and then converted to an estimate of C_{ua} using a carbon fraction constant (Extended Data Table 2).

Root carbon pool (C_{root}) consists of fine root (C_{froot}), intermediate root (C_{iroot}), and coarse root (C_{croot}) pools, with C_{froot} defined as roots with diameter of < 2 mm, C_{iroot} defined as roots with diameter of $2 - 3$ mm, and the remaining roots defined as C_{croot} (Figure S6). The C_{root} pool includes roots of both overstorey and understorey vegetation. Total root biomass (B_{root}) was estimated based on an allometric relationship with stand basal area (derived from DBH) derived for Australian forest species⁴⁴, as follows: $\ln(B_{\text{root}}) = 0.787 \times \ln(\text{DBH}) + 1.218$.

Standing intermediate root (2-3 mm in diameter) and fine root biomass (< 2 mm in diameter) were sampled in four subplots per plot at two depths (0 – 10 cm and 10 – 30 cm) in year 2017, whereas only fine root biomass at the same depths with the same number of subplots was repeatedly sampled over the period of 2014-2016²⁹. We estimated a depth-specific relationship between fine root biomass (< 2 mm in diameter) and total root biomass less than 3 mm in diameter based on samples collected in 2017, and calculated the intermediate root biomass for the period of 2014-2016 based on its corresponding fine root biomass. Coarse root biomass was then estimated as the net difference between total allometrically-derived root biomass and that of roots with diameter < 3 mm. The fine, intermediate, and coarse root

biomass were multiplied by the corresponding carbon fraction constants to obtain C_{froot} , C_{iroot} , and C_{croot} , respectively (Extended Data Table 2).

Microbial carbon pool (C_{micr}) was estimated based on fumigation extraction and 0.5 M K_2SO_4 extraction as in Ref. 25 using samples taken at 0-10 cm soil depth over the period of 2012 - 2015. Total organic carbon was determined on a Shimadzu TOC analyzer (TOC-L TNM-L; Shimadzu, Sydney, Australia), which was then multiplied by soil bulk density over the same soil depth to obtain the C_{micr} (Figure S7a).

Mycorrhizal carbon pool (C_{myco}) for the top 10 cm of soil was estimated via measurements of colonization of mycorrhizal in-growth bags, carbon isotopic partitioning, microbial phospholipid fatty acid abundance and C_{micr} . Nine 45 μm nylon mesh bags (4×5 cm) filled with sand, which excluded roots but allowed access of fungi⁴⁵, were buried in November 2014 in each experimental plot and three bags were subsequently collected every four months for one year. Phospholipid-derived fatty acids (PLFA), a proxy for total microbial biomass abundance, were quantified in sand bags and native field soil following the protocol by Ref 46. $\delta^{13}\text{C}$ values of ground subsamples of this sand, native soil carbon, and aboveground plant tissue (leaves of Eucalypts in April 2014) were used to estimate the fraction of the accumulated carbon in sand bags that was derived from plant carbon using isotopic mass balance. Due to the exclusion of roots, plant-derived carbon in bags can be attributed to mycorrhiza. This plant-derived unitless fraction was then multiplied by the total concentration of PLFA in sand bags to obtain the amount of the total PLFA contributed by mycorrhiza ($\mu\text{g PLFA} / \text{g sand}$). To scale this to native soil PLFA concentrations we then calculated the ratio between mycorrhizal PLFA in sand bags to total PLFA in soil

(representing the total microbial pool). Subsequently, to estimate C_{myco} , this ratio was multiplied by the C_{micr} in each plot (Figure S7b).

Leaf litter carbon pool (C_{lit}) was estimated based on leaf litter decomposition rate and leaf litterfall data collected by litter baskets (Figure S8)²⁴. Leaf litter decomposition rates were estimated over 24 months using litter bags. Briefly, 2 g air-dried *Eucalyptus* litter was added to 10 × 15 cm litter bags with a 2-mm mesh size. Twelve litter bags were randomly allocated to 4 subplots within each treatment plot, and two litter bags were collected at 3, 6, 9, 12, 18 and 24 months to calculate mass loss over time (mass loss was averaged across the two replicates from each subplot). A leaf litter exponential decay function was estimated for each plot, based on data collected over this 24-month period. Leaf litterfall was estimated from monthly collections of material from circular fine-mesh traps (each 0.2 m²) at eight random locations for each plot. We then applied the exponential decay function with litterfall biomass to obtain C_{lit} , assuming a carbon fraction constant (Extended Data Table 2).

Insect carbon pool (C_{ins}) was estimated based on two different sampling techniques, with aerial insects partially estimated based on monthly dead insect data collected from circular fine-mesh traps of 0.2 m² at eight random locations for each plot⁴⁷, and understory insects estimated based on vacuum suction sampling from two locations for each plot⁴⁸. The insect biomass estimated based on these two sampling techniques may be a conservative estimate (the frass produced would suggest presence of a larger insect biomass⁴⁹); nevertheless, they provided a direct estimate based on data collected *in situ*. The vacuum suction method collected invertebrates from understorey vegetation in two 1 × 1 m subplots using a petrol-powered ‘G-Vac’ vacuum device run on full-throttle for 20 s, for a total of five sampling campaigns. Trapping locations were randomly chosen and fixed between sampling

campaigns. All invertebrates were sorted from debris, dried to constant weight at 60 °C and weighed on a microbalance with a precision of 1 µg. We assumed that vacuum samples as well as fine-mesh trap samples represent point estimates of invertebrate abundance. Then, the total biomass of sampled invertebrates was summed across sampling methods within each plot. A constant carbon fraction based on Ref 50 (Extended Data Table 2) was used to convert biomass into C_{ins} pool (Figure S9).

Ecosystem carbon uptake fluxes

Overstorey gross primary production (GPP_O) for each plot was provided by a stand-level model simulation (MAESPA), forced by hourly meteorological data, daily plot-specific leaf area index and leaf-scale treatment-specific photosynthetic parameters measured at the site (Figure S10a)^{7,21}. In short, MAESPA was used as a tool to up-scale leaf-level gas exchange measurements to the whole canopy. In MAESPA, each plot consists of individual tree crowns that are located and parameterized with measured coordinates, crown size, and LAI. Each crown is divided into six layers, with leaf area uniformly distributed in each layer. Within each layer, the model simulates twelve grid points. The incident radiation on the sunlit and shaded leaf area at each grid point is calculated considering shading from upper crown and surrounding trees, solar angle (zenith and azimuth), and light source (diffuse or direct). Incident radiation is then used to calculate gas exchange using a Farquhar⁵¹ formulation for photosynthesis and a Medlyn formulation⁵² for stomatal conductance. The model was parameterized with treatment-specific leaf gas exchange measurements made *in situ*^{7,53}. Leaf respiration and its temperature dependence were also quantified using data collected on site, then up-scaled to the canopy using MAESPA. The performance of the model was evaluated by comparing the simulated transpiration flux to sap flow data⁵⁴.

Similarly, **understorey GPP (GPP_u)** (Figure S10b) was simulated using MAESPA with photosynthetic parameters taken for the dominant grass *Microlaena stipoides*⁴¹. The parameterization of understory vegetation is different from that of the canopy. In each plot, the understory was assumed to form a single crown covering the whole plot (i.e., a circle with 12.5 m radius) at a height of 1.5 m. The LAI of the understory was estimated using phenology camera digital photographs taken at four permanent understorey vegetation monitoring subplots in each plot⁴³. The average green pixel content was calculated from three photos in each subplot, and assumed to be the same as the fraction of absorbed PAR. We then assumed a light extinction coefficient of 0.5 in Beers' Law and calculated understorey LAI. Before 2014 there were 3 campaigns per year while from 2014 the cameras were automated, and we used the fortnightly averages. Leaf gas exchange parameters were obtained from Ref 41 and covered four to six campaigns per year from 2013 to 2016. We estimated a one-time g_l parameter⁵² for all plots and time, and assumed constant carboxylation rate (V_{cmax}) and electron transport rate (J_{max}) values at 25 °C across plots. Basal leaf respiration rate and the temperature dependence of photosynthesis and respiration were assumed to be the same as those for the canopy. The understory simulation was conducted separately from the canopy, with canopy LAI from Ref 24 included to account for the shading from the canopy, branches and stems on the understory.

For the **methane net flux (CH_4)**, air samples were collected following the closed-chamber method (or Non-Flow-Through Non-Steady-State [NFT-NSS] method). Seven replicated chambers were available for each plot. Headspace samples were collected monthly, over a period of one hour and analyzed by gas chromatography. Fluxes were estimated by a mixture of linear and quadratic regressions (depending on goodness-of-fit), assuming a constant air pressure of one atmosphere and correcting the air temperature inside the chambers for each

air sample⁵⁵. The CH₄ fluxes are net fluxes, which represent the sum of: 1) CH₄ efflux (emissions from the soil into the atmosphere); 2) CH₄ influx (uptake from the atmosphere into soil). Here, the annual net CH₄ flux was an ecosystem influx and was presented as positive values (Figure S11a).

Production fluxes

Plant **net primary production (NPP)** is the sum of overstorey leaf (NPP_{ol}), stem (NPP_{stem}), fine root (NPP_{froot}), intermediate root (NPP_{iroot}), coarse root (NPP_{croot}), other (including twigs, barks, and seeds; NPP_{other}), understorey aboveground (NPP_{ua}), and consumption of overstorey leaf by insect herbivores (NPP_{ins}). NPP_{ol} and NPP_{other} were estimated based on monthly litter data collected from circular fine-mesh traps of 0.2 m² at eight random locations for each plot (Figure S12). Litter was sorted into leaf, twigs, bark, and seeds, dried to constant mass at 40 °C and weighed. A subsample was reweighed when dried to constant mass at 70 °C and a small moisture correction⁷ was applied to the leaf component of the whole dataset. NPP_{ol} was computed as the sum of annual leaf litter, which excluded leaf consumption by insects. For twigs, we assumed strictly annual turnover across the years. NPP_{stem} (Figure S13) and NPP_{croot} (Figure S14) were estimated based on annual incremental change of stem biomass and coarse root biomass, respectively. NPP_{froot} was estimated based on samples collected from in-growth cores at four different locations per plot (Figure S14). NPP_{iroot} was estimated based on a global mean coarse root turnover rate (0.3605 yr⁻¹) for evergreen broadleaf forests⁵⁶, and the C_{iroot} pool in our dataset (Figure S14).

NPP_{ua} was estimated based on biomass clippings taken between 2015 - 2017, assuming one understorey turnover per harvest interval (Figure S15). We used a clip-strip method of biomass harvest as has been applied previously at the BioCON experiment⁵⁷. Specifically,

four narrow strips, each with a size of 1 m × 0.1 m, were situated in each of the experimental plots at least 2 m away from the vertical pipes for FACE, while avoiding the understory shrubs. The understory herbaceous species were clipped approximately 1 cm above soil level. The total mass per harvest represents the total production. Biomass samples were oven dried for two days at 60 °C, and converted into carbon mass by applying a constant fraction (Extended Data Table 2).

NPP lost to overstorey leaf consumption by insect herbivores (NPP_{ins}) was estimated based on insect frass data (Frass) collected from the circular fine-mesh traps, and a relationship between frass mass and insect-consumed leaf mass derived based on multiple *Eucalyptus* tree species at different CO₂ concentrations (Figure S16a)^{58,59}. Frass was estimated based on annual collection of frass biomass collected from the circular fine-mesh litter traps with their associated carbon content (Extended Data Table 2; Figure S16c).

Outfluxes

Leaching lost as **dissolved organic carbon (DOC)** from soils was estimated based on concentrations of DOC in soil solutions, provided by water suction lysimeter measurements²⁸. Lysimeters were installed to two depths (0 - 15 cm and 35 - 75 cm, which is immediately above the impermeable layer). Here we assumed that DOC reaching deeper depth is lost from the system at a rate of 20 ml m⁻² d⁻¹, which is an estimate of the daily drainage rate at the site (Figure S11b).

Plant autotrophic respiration (R_a) consists of overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understory aboveground (R_{ua}) (Figure S17), and growth respiration (R_{grow}) (Figure S18). R_{ol}

and R_{ua} were based on MAESPA simulation (Figure S17a, c), as described in the respective GPP sections. R_{grow} was estimated by taking a constant fraction of 30% of total NPP as measured directly on *E. tereticornis* trees⁶⁰.

R_{stem} was estimated from measurements of stem CO_2 efflux performed in three dominant trees per plot (Figure S17b). Collars were horizontally attached to the stem at an approximate height of 0.75 m, and R_{stem} ($nmol\ CO_2\ m^{-2}\ s^{-1}$) was measured with a portable infrared gas analyzer coupled to a soil respiration chamber adapted for this purpose⁶¹. Measurement campaigns were performed every one or two months from December 2017 to October 2018, and the relationship between R_{stem} and air temperature (T_{air}) was used to extrapolate R_{stem} across the surveyed period, following $R_{stem} = 0.1866 \times 2.84^{T_{air}/10}$ ($r^2 = 0.42$, $p < 0.0001$). R_{stem} was then upscaled to the stand level considering the ratio of stem axial surface per unit of soil surface measured per plot. Stem surface area was inferred from the measured tree diameter based on dendrometer, and a relationship between diameter and stem surface area estimated from the Terrestrial Laser Scanning (TLS) data. Stem surface area and diameter in the TLS data was estimated through quantitative structure models presented in Ref. 62 and 63. TLS data were acquired with a RIEGL VC-400 terrestrial laser scanner (RIEGL Laser Measurement Systems GmbH). Stem surface area was derived from the TLS data following a two-step approach: (i) manually extracting single trees from the registered TLS point cloud; and (ii) deriving parameters for an extracted single tree. Once a tree is extracted from the point cloud, the next step was to strip off the leaves, and segment the point cloud into stem and branches. Finally, the surface of the segments was reconstructed with geometric primitives (cylinders). The method used a cover set approach, where the point cloud was partitioned into small subsets, which correspond to small connected patches in the tree surface.

518

519 R_{root} was partitioned into fine root (R_{froot}), intermediate root (R_{iroot}), and coarse root (R_{croot})
520 respiration (Figure S17d). Mass-based rates of fine root and intermediate root respiration
521 ($\text{nmol CO}_2 \text{ DM g}^{-1} \text{ s}^{-1}$) were measured for detached roots sampled by soil cores at 10 cm soil
522 depth at four subplots per plot with a portable infrared gas analyzer coupled to a small root
523 chamber. Measurement campaigns were performed every one or two months from November
524 2018 to July 2019. The relationship between root respiration and soil temperature (T_{soil}) at 10
525 cm soil depth was used to extrapolate the corresponding root respiration rates across the
526 surveyed period, following the equations: $R_{\text{froot}} = 1.138 \times 1.614^{0.0479 \times T_{\text{soil}}}$ ($r^2 = 0.36$, $p <$
527 0.0001 , $\text{RMSE} = 1.054$), and $R_{\text{iroot}} = 0.9764 \times 1.586^{0.0641 \times T_{\text{soil}}}$ ($r^2 = 0.52$, $p < 0.0001$, $\text{RMSE} =$
528 0.597). The mass-based rate of coarse root respiration was assumed to be the same as the
529 mass-based rate of stem respiration. R_{froot} , R_{iroot} and R_{croot} were then upscaled to the stand
530 level to obtain R_{root} with fine root, intermediate root, and coarse root biomass, respectively.

531

532 **Carbon efflux due to insect respiration (R_{ins})** was estimated as the net difference between
533 NPP_{ins} and Frass, assuming no net change in insect biomass (Figure S16b).

534

535 **Soil respiration (R_{soil}):** The rate of soil CO_2 efflux was measured at eight locations within
536 each plot, where a permanent PVC collar inserted into the soil was co-located with soil TDR
537 probes for continuous measurements of soil temperature (5-cm-depth) and volumetric water
538 content (0 to 21-cm-depth; CS650-L; Campbell Scientific, Logan, UT, USA). R_{soil} was
539 measured manually at all collar locations every 2-3 weeks, in addition to 30-minute
540 measurements using automated chambers (Li-8100-103; Licor) at one location within each
541 plot, resulting in $>300,000$ observations over the study period²⁶. These data were used to
542 parameterize a semi-mechanistic model of R_{soil} , in which R_{soil} was predicted based on

measurements of soil properties, soil physics, and measured soil temperature and volumetric water content⁶⁴. This model successfully recreated the observed fluxes (r^2 between predicted and observed survey R_{soil} was 0.65)²⁶. Annual sums of R_{soil} were derived by summing the averaged daily fluxes over eight locations within each plot, where daily fluxes at each location were predicted based on the semi-mechanistic model and daily soil temperature and volumetric water content data taken adjacent to each measurement collar. Soil heterotrophic respiration (R_{hetero}) was taken as the net difference between R_{soil} and R_{root} (Figure S19). Total ecosystem respiration (R) was calculated as the sum of R_a , R_{hetero} , R_{ins} , and VC.

Volatile carbon (VC; Figure S20) flux as isoprene (C_5H_8) and monoterpenes was estimated using the Model of Emissions of Gases and Aerosols from Nature (MEGAN)⁶⁵. Isoprene represents over half of all volatile organic carbon species emitted by vegetation globally, and is the dominant source of VC emission at our site. A MEGAN box-model was built from the version used in Ref. 66, centered on the EucFACE facility to calculate hourly emissions of isoprene across the period 2013-2016 for all six plots:

$$VC = EF \times LAI \times \gamma$$

Where EF is the compound-specific basal emission factor, γ is the emission activity factor, accounting for changes in the emission response due to light, temperature, leaf age and soil moisture. The MEGAN simulations were driven by daily input data of LAI, soil moisture, and hourly input data of photosynthetic active radiation, temperature, atmospheric pressure, wind speed and relative humidity.

The isoprene EFs for ambient and elevated CO_2 plots were derived from in-line photosynthetic gas-exchange measurements coupled with simultaneous volatile isoprenoid sampling. The isoprene was collected onto sterile stainless steel thermal desorption tubes at

the same time as gas exchange was measured, and these were capped and later thermally desorbed for off-line volatile analysis in the laboratory using a Shimadzu 2010 Plus GC-MS system connected to a Shimadzu TD20 automated cartridge desorber. The sampling and GC-MS analysis methodology is described in detail in Ref 67. The chromatographic peaks were identified by comparing them to an isoprene standard and reference mass spectra in the NIST Mass Spectral Library (<https://www.nist.gov/srd>). Monoterpene emissions were sampled during February 2018 using a push-pull headspace technique⁶⁸ from enclosed branches containing approximately 10 leaves and trapped on adsorbent cartridges (150 mg Tenax TA and 200 mg Carbograph 1TD, Markets International Limited, United Kingdom) at an outflow rate of 200 ml min⁻¹ for 15 min. Before each measurement, the sampling system was equilibrated for 15 min at an inflow rate of 1000 ml min⁻¹. Monoterpenes were analyzed by gas chromatography-mass spectrometry (R7890A Series GC coupled with a 5975C inert MSD/DS Performance Turbo EI System, Agilent Technologies, Inc., Santa Clara, CA, USA), as described by Ref 69. The obtained chromatograms were deconvoluted, analyzed and data retrieved using the software PARADISE⁷⁰ version 3.88. Identification of compounds was performed using analytical standards and according to their mass spectra in the NIST11 library. Pure analytical standards were used for quantification. The box-model produced isoprene and monoterpenes were converted to carbon content using the respective molecular mass ratios.

Net Ecosystem Production

Net ecosystem production (NEP) was estimated based on three different methods that estimated NEP in relatively independent ways (Figure 3), similar to Ref 71. The first method considered NEP as the difference between total ecosystem influx and total ecosystem outflux (i.e. In - Out), which relied on both process-based modeling and empirical upscaling of

respiratory fluxes collected from the field. The second method considered NEP as NPP minus R_{hetero} (i.e. $\text{NPP} - R_{\text{hetero}}$), with NPP relying mostly on litter-based production estimates, and R_{hetero} relying on R_{soil} and R_{root} estimates. The third method considers NEP as the sum of changes in carbon pools over time in the ecosystem (i.e. ΔC_{pools}), which was mostly determined by biomass estimates. Equations for each method are provided below:

Method	NEP =
In - Out	$\text{GPP}_{\text{o}} + \text{GPP}_{\text{u}} + \text{CH}_4 - R_{\text{ol}} - R_{\text{stem}} - R_{\text{soil}} - R_{\text{ua}} - R_{\text{ins}} - \text{DOC} - \text{VC} - R_{\text{grow}}$
$\text{NPP} - R_{\text{hetero}}$	$\text{NPP}_{\text{ol}} + \text{NPP}_{\text{stem}} + \text{NPP}_{\text{froot}} + \text{NPP}_{\text{iroot}} + \text{NPP}_{\text{croot}} + \text{NPP}_{\text{other}} + \text{NPP}_{\text{ua}} + \text{NPP}_{\text{ins}} - R_{\text{hetero}}$
ΔC_{pools}	$\Delta C_{\text{soil}} + \Delta C_{\text{ol}} + \Delta C_{\text{stem}} + \Delta C_{\text{croot}} + \Delta C_{\text{froot}} + \Delta C_{\text{iroot}} + \Delta C_{\text{ua}} + \Delta C_{\text{lit}} + \Delta C_{\text{ins}} + \Delta C_{\text{micr}} + \Delta C_{\text{myco}}$

Carbon budget evaluation

We evaluated the mass balance of our estimated ecosystem carbon budget in two ways. Firstly, we compared model simulated GPP with the aggregated sum of NPP and R_{a} (Extended Data Figure 3a, b). GPP was simulated by a stand-level ecophysiological model, driven by hourly meteorological data and parameterized with site-specific ecological data²⁰. This GPP should equal to the aggregation of NPP ($\text{NPP}_{\text{ol}} + \text{NPP}_{\text{stem}} + \text{NPP}_{\text{froot}} + \text{NPP}_{\text{iroot}} + \text{NPP}_{\text{croot}} + \text{NPP}_{\text{other}} + \text{NPP}_{\text{ua}} + \text{NPP}_{\text{ins}}$) and R_{a} fluxes ($R_{\text{ol}} + R_{\text{stem}} + R_{\text{root}} + R_{\text{ua}} + R_{\text{grow}}$), which were mostly extrapolated based on field data. Secondly, R_{soil} estimated based on soil collar flux measurements²⁴ was evaluated against the sum of litterfall and R_{root} (Extended Data Figure 3c, d), assuming minimal changes in soil carbon stock (as change over this short period of time is beyond the detection limit in a complex and slow-growing mature forest ecosystem like EucFACE). Here, litterfall was the sum of $\text{NPP}_{\text{ol}} + \text{NPP}_{\text{froot}} + \text{NPP}_{\text{iroot}} +$

NPP_{other} + NPP_{ua} + Frass, and R_{root} was extrapolated based on root biomass and temperature functions.

Statistical analyses

We performed linear mixed-model analysis using the “lmer” function within the “lme4” package⁷² in software R⁷³ to determine the CO₂ treatment effect on all reported variables. All fluxes were reported at an annual rate (g C m⁻² yr⁻¹). In our model, date and CO₂ treatment were considered as fixed factors, plot as a random factor, and plot-specific pre-treatment LAI (i.e. 4-month average LAI before full CO₂ treatment was switched on) as a covariate to account for pre-treatment differences among treatment plots. Normalizing all response variables with a covariate that integrates light, water and nutrient constraints helps to isolate the CO₂ effect²³, as has been done previously at the site²⁴ and elsewhere^{8,23}. Confidence intervals for the CO₂ effect size of individual variables were reported using the function “confint”, which applies quantile functions for the t-distribution after model fitting. Confidence intervals for the predicted flux and pool were reported as the standard deviation of the plot-specific totals (n = 3). Similarly, confidence intervals for the aggregated fluxes (e.g. NPP) were reported by summing individual component fluxes that constitutes the aggregated flux for each plot and computing the standard deviations across plots (n = 3). Finally, confidence intervals for the CO₂ effect size (SD_{agg}) of some aggregated fluxes (e.g. NPP) were calculated by pooling the standard deviations of the aggregated fluxes for ambient (SD_{amb}) and elevated CO₂ treatment (SD_{ele}), following:

$$SD_{agg} = \sqrt{\frac{SD_{amb}^2 + SD_{ele}^2}{2}}$$

Uncertainty analysis

We applied a Markov Chain Monte Carlo (MCMC) data assimilation algorithm to a simplified carbon cycle framework to make inference of the uncertainties around the fate of carbon in our carbon budget. We simplified our carbon budget into eight pools (Extended Data Figure 7), namely, leaf (C'_{leaf} , which includes overstorey and understorey), wood (C'_{wood} , which includes stem and coarse root), root (C'_{root} , which includes fine root and intermediate root), aboveground litter (C'_{aglit}), belowground litter (C'_{bglit}), mycorrhizae (C'_{myco}), microbe (C'_{micr}), and soil (C'_{soil}). Here, C'_{aglit} and C'_{bglit} were assumed unknowns and inferred from the analysis. Net primary production (NPP) was calculated as the difference of gross primary production (GPP) and autotrophic respiration (R_a). NPP was then allocated into the four plant carbon pools (C'_{leaf} , C'_{wood} , C'_{root} , and C'_{myco}), with the respective fitted allocation coefficients (a_{leaf} , a_{wood} , a_{root} , and a_{myco}) being inferred. It has been shown that plant carbon allocation to mycorrhizal fungi may be an important flux in forest carbon budget calculation⁷⁴. Turnover rates of C'_{leaf} , C'_{root} , C'_{myco} , C'_{aglit} , C'_{bglit} , C'_{micr} and C'_{soil} were represented by the corresponding turnover coefficients (τ_{leaf} , τ_{wood} , τ_{root} , τ_{myco} , τ_{aglit} , τ_{bglit} , τ_{micr} , τ_{soil}), all of which were assumed unknowns except τ_{wood} (estimated based on litter basket data of twigs, barks and seeds) and τ_{aglit} (estimated from the leaf litter decomposition data). For carbon leaving from C'_{aglit} , C'_{bglit} and C'_{micr} , we inferred the corresponding fractional coefficient that determines the fraction of carbon entering into the next pool (f'_{aglit} , f'_{bglit} , and f'_{micr}), and assumed the remainder to be respired as part of R_{hetero} . The turnover of soil carbon (i.e. τ_{soil}) also contributed to R_{hetero} . In total, we fitted 2 pools, 4 allocation coefficients, 6 turnover rates, and 3 fractional coefficients using the MCMC algorithm.

We used plot-level estimates of GPP, R_a , R_{hetero} , carbon pools and changes in pools to constrain the MCMC fitting. We assumed uniform parameter distributions and a burn-in

coefficient of 10%. Chain lengths were set at 200,000 for the ambient CO₂ plots and 500,000 for the elevated plots. The longer chain length for the elevated plots was due to the smaller proposal step size for these plots to meet an acceptance rate of around 20%. We reported the means and standard deviation of the estimated parameters at the treatment level in Extended Data Figure 7.

Data statement

Data will be available via Figshare (DOI: 10.6084/m9.figshare.11634315) with the publication of the manuscript. Code to process the data is available via GitHub (https://github.com/mingkaijiang/EucFACE_Carbon_Budget/releases/tag/V20200120).

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Acknowledgements

EucFACE was built as an initiative of the Australian Government as part of the Nation-building Economic Stimulus Package, and is supported by the Australian Commonwealth in collaboration with Western Sydney University. We acknowledge the technical support by V. Kumar, C. McNamara and S. Wohl, and the team of people who have assisted with data collection. The *Eucalyptus* tree vector in Figure 1 is from Heydon, L. *Eucalyptus* spp. Integration and Application Network, University of Maryland Center for Environmental Science (ian.umces.edu/imagelibrary/). This work was partially supported by the following grants from the Australian Research Council (ARC): DP130102501 (to JRP and ICA), DP170104634 (to BKS and PBR), DP110105102 and DP160102452 (to DSE). MGDK acknowledges funding from the ARC Centre of Excellence for Climate Extremes (CE170100023), the ARC Discovery Grant (DP190101823) and support from the NSW Research Attraction and Acceleration Program. RLS received funding from Research Foundation Flanders and the European Union's Horizon 2020 research and innovation programme under the Marie Skłodowska Curie grant agreement no. 665501. RO-H. is financially supported by a Ramón y Cajal Fellowship from MICIU (RYC-2017-22032). EHJN and BMDS received funding from VILLUM Center for Plant Plasticity (VKR023054), VILLUM Young Investor Program fellowship (VKR013167), and a Danish Independent Research Council Sapere Aude Research Talent Post-Doctoral Stipend (6111-00379B). ÜN and AK have been supported by the European Commission through the European Regional Fund (Center of Excellence EcolChange).

905 **Author contributions**

906 MJ, BEM, RAD and JED designed the synthesis, compiled the data, and performed the
907 analyses. MJ, BEM, JED, RAD, ICA, CVMB, MMB, YC, LC-G, LC, KYC, BMDS, SLF,
908 ANG, TEG, SH, SNJ, AK, CAM, KM, BDM, LN, EHJN, UNN, ÜN, NJN, RO-H, VSP, EP,
909 JP, JP, JRP, SAP, PBR, AAR, MR, RR, PR, RLS, BKS, BS, MGT, JKMW, AW-K, JY and
910 DSE collected data and contributed to data analyses. MJ performed data assimilation analysis,
911 with contributions from MGDK and BEM. JY and BEM performed the MAESPA model
912 simulations, with contributions from MGDK and RAD. JED and AAR performed soil
913 respiration gap-filling and modelling. KME performed the MEGAN model simulation. MJ
914 and LC-G conceptualized Figure 1, and LC-G implemented the graphic design. MJ wrote the
915 initial manuscript, with significant input from BEM, JED, BS, PBR, SZ, MGDK, MGT and
916 DSE. All authors edited and approved the manuscript.

917

918 **Competing financial interests**

919 None declared.

920

921 **Materials and Correspondence**

922 Correspondence should be directed to MJ (m.jiang@westernsydney.edu.au) and BEM
923 (b.medlyn@westernsydney.edu.au).

Figure legend

Figure 1. A comprehensive carbon budget under ambient and elevated CO₂ treatment in a mature forest ecosystem.

Diamond boxes are gross primary production for overstorey (GPP_o) and understorey (GPP_u), respectively. Squared boxes are average carbon stocks over the experimental period (C_{pools}, g C m⁻²), including overstorey leaf (C_{ol}), stem (C_{stem}), coarse root (C_{croot}), fine root (C_{froot}), intermediate root (C_{iroot}), understorey aboveground (C_{ua}), leaf litter (C_{lit}), soil (C_{soil}), microbe (C_{micr}), aboveground insect (C_{ins}), and mycorrhizae (C_{myco}). Unboxed variables are carbon fluxes (g C m⁻² yr⁻¹), including net primary production of overstorey leaf (NPP_{ol}), stem (NPP_{stem}), coarse root (NPP_{croot}), fine root (NPP_{froot}), intermediate root (NPP_{iroot}), and understorey aboveground (NPP_{ua}), overstorey leaf consumption by insects (NPP_{ins}), respiration fluxes of overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understorey aboveground (R_{ua}), growth (R_{grow}), insect (R_{ins}), heterotroph (R_{hetero}), and soil (R_{soil}), and volatile carbon emission (VC), frass production (Frass), dissolved organic carbon (DOC), and soil methane net uptake (CH₄). Solid arrow lines are fluxes entering a pool, dotted arrow lines are fluxes leaving a pool. The changes in each carbon pool over time (ΔC_{pools} , g C m⁻² yr⁻¹) are reported in Extended Data Figure 6. Blue italic values are means \pm one standard deviation of the ambient CO₂ treatment (n=3), whereas red values are means \pm one standard deviation of the elevated CO₂ treatment (n=3). All values are normalized by a linear mixed-model with plot-specific pre-treatment leaf area index as a covariate to account for pre-existing differences. A summary of variable definitions and data availability is provided in Extended Data Table 1.

Figure 2. The fate of additional carbon fixed under elevated CO₂ (eCO₂) in a mature forest ecosystem. **a)** Column “GPP” represents the total eCO₂-induced increases in overstorey and understorey gross primary production (GPP_o and GPP_u, respectively), “NPP + R_a” represents the sum of net primary production and autotrophic respiration response, “R + ΔC_{pools}” represents the sum of ecosystem respiration and change in carbon storage response. **b)** The relative contributions of individual NPP fluxes to the aggregated NPP response to eCO₂, including NPP responses of overstorey leaf (NPP_{ol}), twigs, barks and seeds (NPP_{other}), fine root (NPP_{fr}), and understorey aboveground (NPP_{ua}); **c)** The relative contributions of individual respiratory fluxes to the aggregated R response to eCO₂, including respiration responses of stem (R_{stem}), root (R_{root}), understorey aboveground (R_{ua}), growth (R_{grow}), and soil heterotroph (R_{hetero}); and **d)** The relative contributions of individual change in carbon storage to the aggregated ΔC_{pools} response to eCO₂, including changes in pool of stem (ΔC_{stem}), understorey aboveground (ΔC_{ua}), fine root (ΔC_{fr}), leaf litter (ΔC_{lit}), and soil (ΔC_{soil}). Variables with an absolute mean CO₂ effect of < 5 g C m⁻² yr⁻¹ are not reported in the bar chart for better visual clarification. Individual CO₂ responses are reported in Extended Data Figure 5. Each color represents the CO₂ response of a flux variable, the point indicates the net sum of all variables for a column, and the grey error bar represents one standard deviation of the estimated column sum at the plot-level (see Methods). The CO₂ effect is estimated using a linear mixed-model analysis with plot-specific pre-treatment leaf area index as a covariate to account for pre-existing differences (see Methods). The non-normalized response is provided in Extended Data Figure 4, which generally agrees with findings present in this figure, but with larger uncertainty.

Figure 3. Estimates of net ecosystem production (NEP) under ambient and elevated CO₂ treatment at EucFACE. Positive values indicate ecosystem net carbon uptake by the ecosystem. “In - Out” calculates NEP based on the difference between total influxes and total outfluxes. “NPP - R_{hetero}” calculates NEP based on the difference between net primary production (NPP) and heterotrophic respiration (R_{hetero}). “ΔC_{pools}” derives NEP based on incremental changes in all ecosystem carbon pools. Colored bars indicate treatment means based on each method (n=3), with blue representing ambient and red representing elevated CO₂ treatment. Individual dots are plot-level NEP, derived based on different methods (see Methods). Values are normalized by a linear mixed-model with plot-specific pre-treatment leaf area index as a covariate to account for pre-existing differences. Horizontal dotted line indicates NEP equals zero. The inset figure includes an inferred production allocation flux to mycorrhizal fungi (NPP_{myco}) based on data assimilation (Methods), which affected NEP estimates based on the NPP – R_{hetero} method only.

981 **Extended Data Table 1. Definition and data availability of variables.** Data availability
982 includes start and end year of data included in this study. Time points indicate the number of
983 data collections over the available data period. Within plot sub-replicate indicate the number
984 of replicates within each treatment plot. The detailed methods for estimating each variable is
985 provided in the Method section.

Variable		Data coverage			
Name	Symbol	Start year	End year	Time points	Within plot sub-replicate (plot ⁻¹)
Specific Leaf Area	SLA	2013	2016	50	3
Leaf Area Index	LAI	2012	2016	303	1
Soil bulk density	BK	2017	2017	2	3
Diameter at breast height	DBH	2013	2016	4	Individual tree
Overstorey leaf pool	C _{ol}	2012	2016	303	1
Understorey aboveground pool	C _{ua}	2015	2016	16	4
Overstorey stem C pool	C _{stem}	2013	2016	4	Individual tree
Fine root C pool	C _{froot}	2014	2016	7	4
Intermediate root C pool	C _{iroot}	2014	2016	7	4
Coarse root C pool	C _{croot}	2013	2016	4	Individual tree
Forest floor leaf litter C pool	C _{lit}	2013	2016	46	-
Microbial C pool	C _{micr}	2012	2015	15	4
Soil C pool	C _{soil}	2012	2014	11	4

Mycorrhizal C pool	C_{myco}	2015	2015	3	-
Insect C pool (aerial)	C_{ins}	2013	2016	43	8
Insect C pool (understorey)	C_{ins}	2014	2015	5	2
Overstorey gross primary production	GPP_o	2013	2016	Annual	1
Understorey gross primary production	GPP_u	2013	2016	Annual	1
Overstorey leaf respiration	R_{ol}	2013	2016	Annual	1
Understorey leaf respiration	R_{ua}	2013	2016	Annual	1
Stem respiration	R_{stem}	2012	2016	Daily	3
Root respiration	R_{root}	2012	2015	Daily	-
Methane net flux	CH_4	2013	2016	35	7
Volatile C emission flux	VC	2013	2016	Daily	1
Insect herbivore respiration	R_{ins}	2012	2014	22	-
Dissolved organic C loss flux	DOC	2012	2014	12	4
Soil respiration	R_{soil}	2012	2015	Daily	8
Growth respiration	R_{grow}	2012	2016	Annual	1
Overstorey leaf net primary production	NPP_{ol}	2012	2016	49	8
Stem net primary production	NPP_{stem}	2012	2016	4	Individual tree
Fine root net primary production	NPP_{froot}	2014	2016	6	4
Intermediate root net primary production	NPP_{iroot}	2014	2016	6	4

Coarse root net primary production	NPP_{croot}	2012	2016	4	Individual tree
Other net primary production (sum of twigs, bark, seeds)	NPP_{other}	2012	2016	49	8
Twig net primary production	NPP_{twig}	2012	2016	49	8
Bark net primary production	NPP_{bark}	2012	2016	49	8
Seed net primary production	NPP_{seed}	2012	2016	49	8
Understorey aboveground net primary production	NPP_{ua}	2015	2016	3	4
Frass production	Frass	2012	2014	22	8
Heterotrophic respiration	R_{hetero}	2012	2016	Daily	8
Overstorey leaf insect consumption flux	NPP_{ins}	2012	2014	22	-

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987 **Extended Data Table 2. Carbon (C) fraction used to convert from biomass into C**
988 **content.**

Variable	Symbol	Mean value	Data source
C fraction of overstorey leaf pool	f_{ol}	0.5	EucFACE data
C fraction of understorey aboveground pool	f_{ua}	0.456	EucFACE data
C fraction of stem pool	f_{stem}	0.445 (ambient plots) 0.448 (elevated plots)	EucFACE data
C fraction of coarse root pool	f_{croot}	0.445 (ambient plots) 0.448 (elevated plots)	Assumed the same as f_{stem}
C fraction of fine root pool	f_{froot}	0.40 (ambient plots) 0.42 (elevated plots)	EucFACE data
C fraction of intermediate root pool	F_{iroot}	0.40 (ambient plots) 0.42 (elevated plots)	Assumed the same as f_{froot}
C fraction of overstorey leaf litter pool	f_{lit}	0.5	EucFACE data
C fraction of aboveground insect pool	f_{ins}	0.5	Ref 49
C fraction of frass production	f_{frass}	0.53	EucFACE data
C fraction of microbial pool	f_{micr}	0.534 (ambient plots) 0.493 (elevated plots)	EucFACE data
C fraction of mycorrhizal pool	f_{myco}	0.534 (ambient plots) 0.493 (elevated plots)	Assumed the same as f_{micr}
C fraction of soil pool	f_{soil}	0.016 (ambient plots) 0.017 (elevated plots)	EucFACE data
C fraction of twigs, barks and seeds production	f_{other}	0.5	Assumed

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Extended Data Figure 1. The *Eucalyptus* Free Air Carbon dioxide Enrichment experiment facility (EucFACE). **a)** View of the forest and facility from above (photo credit: David S. Ellsworth), **b)** view of the understorey vegetation and infrastructure inside a plot (photo credit: Mingkai Jiang), and **c)** view from below of the canopy structure and the crane (photo credit: Mingkai Jiang).

Extended Data Figure 2. Mean annual temperature (MAT) and mean annual precipitation (MAP) for major forest biomes and a selected list of tree-based elevated CO₂ experiments. Gridded temperature and precipitation data were obtained from the Climate Research Unit (CRU) monthly dataset at 0.5 resolution⁷⁵. Global biome boundaries and definitions were taken from Ref 76 and were spatially aggregated onto the CRU resolution, following Ref 77. The major forest biomes are defined as: tropical and subtropical moist broadleaf forests; tropical and subtropical dry broadleaf forests; tropical and subtropical coniferous forest; temperate broadleaf and mixed forests; temperate coniferous forests; boreal forests/taiga; and Mediterranean forests, woodlands, and scrub. The list of elevated CO₂ experiments includes 7 Free Air CO₂ Enrichment experiments (FACE) and a Whole-Tree Chamber experiment (WTC), namely: EucFACE, DukeFACE, ORNLFACE, AspenFACE, PopFACE, WebFACE, BiForFACE, and FlakalidenWTC. The site-specific climate, tree age and net primary production (NPP) under ambient CO₂ treatment were collected from Ref 3, 9, 10, 11, 78 and 79. The top inset figure compares global forest NPP against standing age using data collected from Ref 80. We included data with forest age < 500 years, and the NPP reported in Ref 80 included both overstorey and understorey. The bottom inset figure compares soil total nitrogen and labile phosphorus across the eCO₂ experiments. Soil total nitrogen was extracted from Ref 81 using spatial coordinates of each experiment, while soil labile phosphorus was spatially extracted from Ref 82. The two dotted lines indicates N:P ratios of 20:1 and 100:1, respectively.

Extended Data Figure 3. Estimates of (a and b) gross primary production (GPP) and (c and d) soil respiration (R_{soil}) based on different methods for both (a and c) ambient and (b and d) elevated CO_2 treatment at EucFACE. For estimates of GPP, we compared the model simulated total GPP of overstorey and understorey (GPP_o and GPP_u , respectively), with the sum of data-driven estimates of net primary production (NPP) and autotrophic respiration (R_a), which include NPP of overstorey leaf (NPP_{ol}), stem (NPP_{stem}), fine root ($\text{NPP}_{\text{froot}}$), intermediate root ($\text{NPP}_{\text{iroot}}$), coarse root ($\text{NPP}_{\text{croot}}$), twigs, barks and seeds ($\text{NPP}_{\text{other}}$), understorey aboveground (NPP_{ua}), leaf consumption by insects (NPP_{ins}), and respiratory fluxes of overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understorey aboveground (R_{ua}), growth (R_{grow}), and volatile carbon emission (VC). For estimates of R_{soil} , we compared direct estimates of R_{soil} scaled up from soil chamber measurements, with the sum of litterfall and independent estimates of root respiration ($\text{Litter} + R_{\text{root}}$), assuming no net change in soil carbon stock over time. Here litterfall was inferred based on NPP of overstorey leaf (NPP_{ol}), fine root ($\text{NPP}_{\text{froot}}$), intermediate root ($\text{NPP}_{\text{iroot}}$), twigs, barks and seeds ($\text{NPP}_{\text{other}}$), understorey aboveground (NPP_{ua}), and frass production (Frass). These evaluations provide independent mass balance checks of the estimated ecosystem carbon budget. Each color represents a flux variable. Dotted point and vertical line represent treatment mean and standard deviation based on plot-level estimates of the aggregated flux ($n=3$). Values were normalized by a linear mixed-model with pre-treatment leaf area index as a covariate to account for pre-existing differences.

Extended Data Figure 4. The fate of additional carbon fixed under elevated CO₂ (eCO₂) in a mature forest ecosystem (non-normalized analysis case). **a)** Column “GPP” represents the total eCO₂ induced increase in overstorey and understorey gross primary production (GPP_o and GPP_u, respectively), column “NPP + R_a” represents the sum of net primary production and autotrophic respiration eCO₂ response, and column “R + ΔC_{pools}” represents the sum of ecosystem respiration and carbon storage eCO₂ response. **b)** The relative contributions of individual NPP fluxes to the aggregated NPP response to eCO₂, including overstorey leaf (NPP_{ol}), stem (NPP_{stem}), fine root (NPP_{froot}) and understorey aboveground (NPP_{ua}). **c)** The relative contributions of individual respiratory fluxes to the aggregated R response to eCO₂, including overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understorey aboveground (R_{ua}), and heterotroph (R_{hetero}). **d)** The relative contributions of individual change in carbon storage to the aggregated ΔC_{pools} response to eCO₂, including stem (ΔC_{stem}), fine root (ΔC_{froot}), leaf litter (ΔC_{lit}), understorey aboveground (ΔC_{ua}), and soil (ΔC_{soil}). Variables with an average CO₂ effect of < 5 g C m⁻² yr⁻¹ were excluded from the figure for better visual clarification. Each color represents a flux variable, point indicates the net sum of all variables for a column, and the grey confidence interval represents plot-level standard deviation (n=3) of the estimated column sum.

Extended Data Figure 5. CO₂ treatment effect (g C m⁻² yr⁻¹) for all ecosystem fluxes at EucFACE. **a)** The CO₂ response of gross ecosystem carbon uptake, including gross primary production of overstorey (GPP_o) and understorey (GPP_u), and soil methane uptake (CH₄). **b)** The eCO₂ response of annual incremental change in carbon pool (ΔC_{pools}), including overstorey leaf (ΔC_{ol}), stem (ΔC_{stem}), coarse root (ΔC_{croot}), fine root (ΔC_{froot}), intermediate root (ΔC_{iroot}), understorey aboveground (ΔC_{ua}), leaf litter (ΔC_{lit}), soil (ΔC_{soil}), microbe (ΔC_{micr}), aboveground insect (ΔC_{ins}), and mycorrhizae (ΔC_{myco}). **c)** The eCO₂ response of net primary production (NPP), including overstorey leaf (NPP_{ol}), stem (NPP_{stem}), coarse root (NPP_{croot}), fine root (NPP_{froot}), intermediate root (NPP_{iroot}), understorey aboveground (NPP_{ua}), twigs, barks and seeds (NPP_{other}), and leaf insect consumption (NPP_{ins}). **d)** The eCO₂ response of ecosystem respiration (R) and other out-going flux, including respiration fluxes of overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understorey aboveground (R_{ua}), growth (R_{grow}), insect (R_{ins}), heterotroph (R_{hetero}), and soil (R_{soil}), and volatile carbon emission (VC) and dissolved organic carbon leaching (DOC). Dots and grey bars represent means and standard deviations of the CO₂ treatment difference, predicted by a linear mixed-model with plot-specific pre-treatment leaf area index as a covariate. Red dots indicate negative means and blue dots indicate positive means. Dashed lines indicate change of scale along the x-axis.

Extended Data Figure 6. Estimates of incremental change in carbon pool averaged over the experimental period under ambient (aCO₂) and elevated CO₂ (eCO₂) treatment effect at EucFACE (ΔC_{pools} , g C m⁻² yr⁻¹). The ΔC_{pools} variables are overstorey leaf (ΔC_{ol}), stem (ΔC_{stem}), coarse root (ΔC_{croot}), fine root (ΔC_{froot}), intermediate root (ΔC_{iroot}), understorey aboveground (ΔC_{ua}), leaf litter (ΔC_{lit}), soil (ΔC_{soil}), microbe (ΔC_{micr}), aboveground insect (ΔC_{ins}), and mycorrhizae (ΔC_{myco}). Colored bars and black lines represent means and standard deviations for each treatment, with blue represents aCO₂ and red represents eCO₂ treatment. Dashed lines indicate change of scale along the x-axis.

Extended Data Figure 7. Fitted carbon cycle parameters to trace the fate of the additional carbon under elevated CO₂ at EucFACE. Parameters were estimated by Markov Chain Monte Carlo (MCMC) fitting algorithm, assuming a simplified carbon cycle framework based on data collected from EucFACE. Details of the MCMC approach can be found in the Methods. Plot-level gross primary production (GPP), autotrophic respiration (R_a), heterotrophic respiration (R_{hetero}), carbon pools of leaf (C'_{leaf}), wood (C'_{wood}), root (C'_{root}), mycorrhizae (C'_{myco}), microbe (C'_{micr}), and soil (C'_{soil}), and the corresponding change in pools were used to constrain the model fitting. Net primary production (NPP) was derived as the difference of GPP and R_a . Carbon use efficiency (CUE') was calculated as NPP/GPP ; it differs from the value given in the main text owing to the contribution of NPP allocated to mycorrhizae (NPP_{myco}). We fitted two carbon pools (C'_{aglit} and C'_{bglit}), four allocation coefficients (a_{leaf} , a_{wood} , a_{root} , and a_{myco}), six turnover rates (τ_{leaf} , τ_{root} , τ_{myco} , τ_{bglit} , τ_{micr} , and τ_{soil}), and three fractional coefficients (f'_{aglit} , f'_{bglit} , and f'_{micr}) using MCMC algorithm. The fractional coefficients indicate the fraction of carbon leaving one pool that enters the subsequent pool, with the remainder respired as R_{hetero} .

1096

1097 **Extended Data Figure 8. Data-model intercomparison of some key carbon cycle**
1098 **parameters, under ambient (aCO₂) and elevated CO₂ (eCO₂).** Parameters include: **a)**
1099 allocation coefficients to leaf, wood, root and other, **b)** turnover rates of leaf, root,
1100 aboveground litter (Aglit), belowground litter (Bglit), and **c)** turnover rate of soil. Models
1101 include: Community Atmosphere Biosphere Land Exchange (CABL), Community Land
1102 Model 4 (CLM4), Community Land Model with a phosphorus component (CLMP), Generic
1103 Decomposition And Yield (GDAY), Lund-Potsdam-Jena General Ecosystem Simulator
1104 (LPJX), Orchidee-C-N (OCNX), and Sheffield Dynamic Global Vegetation Model (SDVM).
1105 The model output was generated as part of the model ensemble predictions made in advance
1106 of the experiment reported in Ref 17 for EucFACE. Data-based uncertainties were estimated
1107 using the Markov Chain Monte Carlo data assimilation algorithm, with error bars indicating
1108 one standard deviation. Allocation to other in the data refers to the allocation to mycorrhizal
1109 production, whereas it refers to the allocation to reproductive carbon pool in some models.

1110

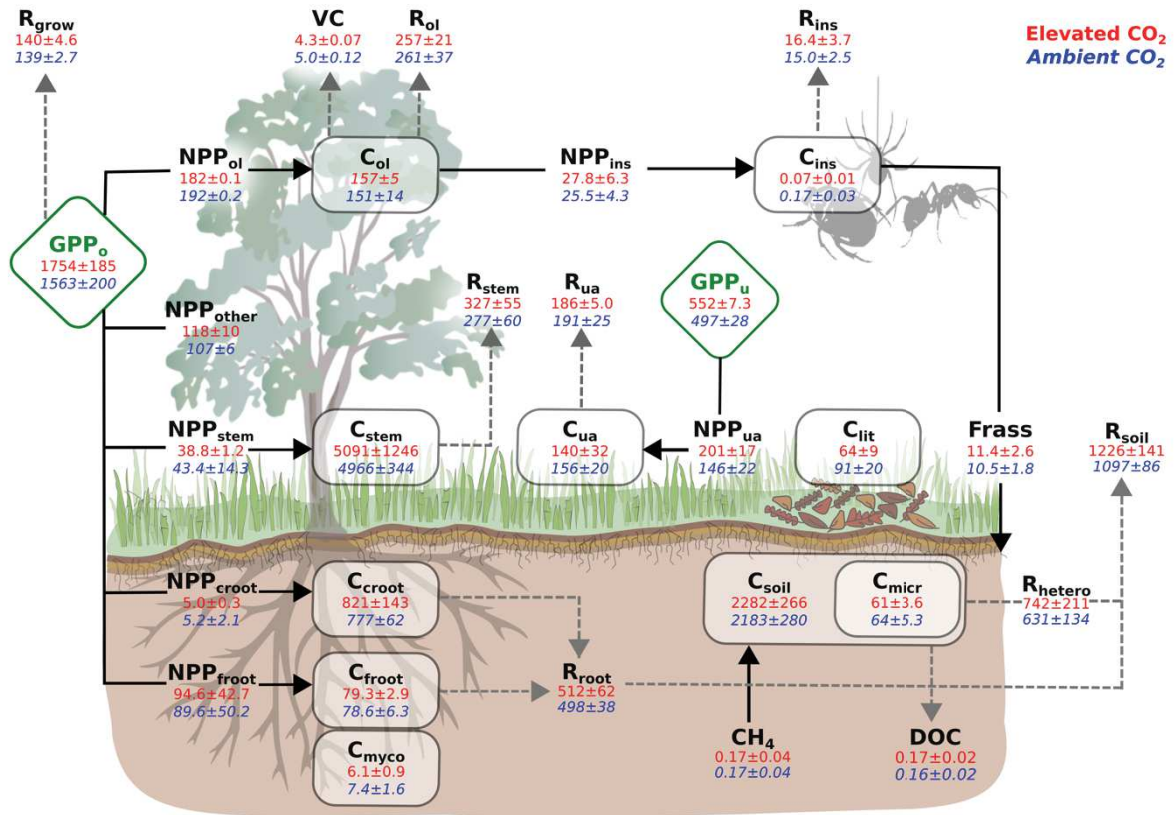
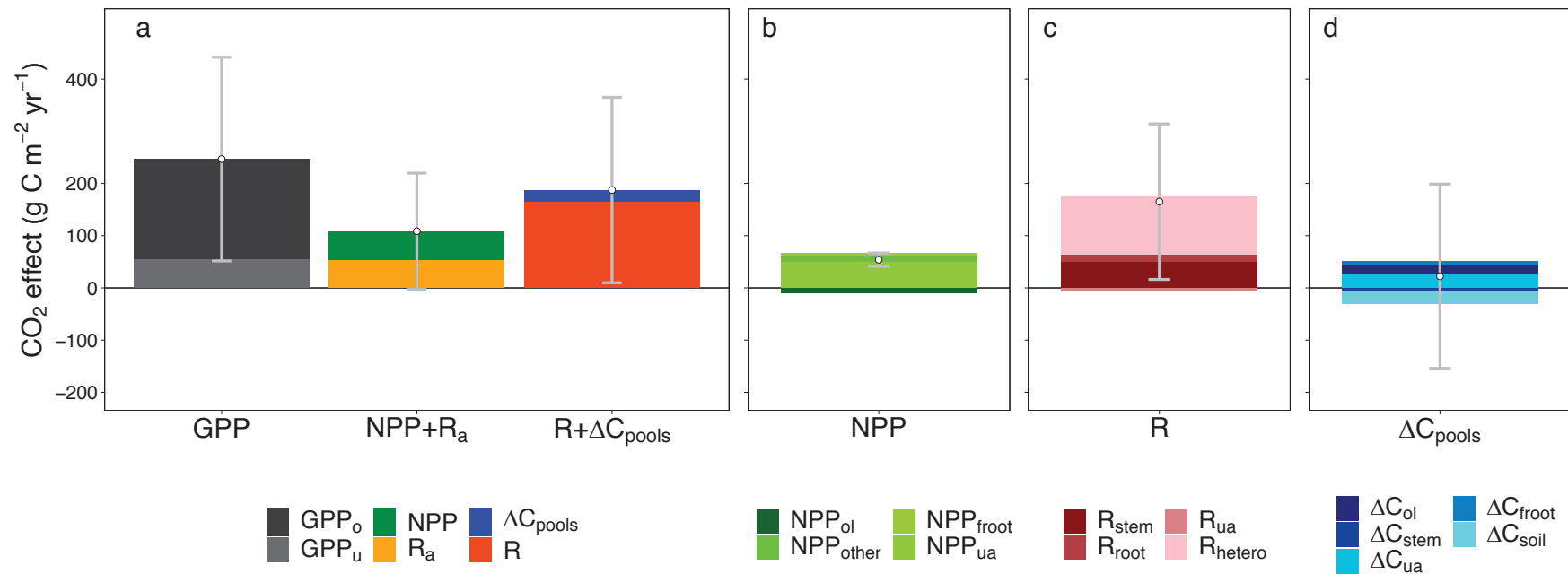


Figure 1. A comprehensive carbon budget under ambient and elevated CO₂ treatment in a mature forest ecosystem. Diamond boxes are gross primary production for overstorey (GPP_o) and understorey (GPP_u), respectively. Squared boxes are carbon stocks (gCm⁻²), including overstorey leaf (C_{ol}), stem (C_{stem}), coarse root (C_{croot}), fineroot (C_{froot}), understorey aboveground (C_{ua}), leaf litter (C_{lit}), soil (C_{soil}), microbe (C_{micr}), aboveground insect (C_{ins}), and mycorrhizae (C_{myco}). Unboxed variables are carbon fluxes (gCm⁻²yr⁻¹), including net primary production of overstorey leaf (NPP_{ol}), stem (NPP_{stem}), coarse root (NPP_{croot}), fineroot (NPP_{froot}), and understorey aboveground (NPP_{ua}), overstorey leaf consumption by insects (NPP_{ins}), respiration fluxes of overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understorey aboveground (R_{ua}), growth (R_{grow}), insect (R_{ins}), heterotroph (R_{hetero}), and soil (R_{soil}), and volatile carbon emission (VC), frass production (Frass), dissolved organic carbon (DOC), and soil methane net

uptake (CH₄). Solid arrow lines are fluxes entering a pool, dotted arrow lines are fluxes leaving a pool. Blue italic values are means \pm one standard deviation of the ambient CO₂ treatment (n=3), whereas red values are means \pm one standard deviation of the elevated CO₂ treatment (n=3). All values are normalized by a linear mixed-model with plot-specific pre-treatment leaf area index as a covariate to account for pre-existing differences. Summary of variable definitions and data availability is provided in Extended Data Table 1.

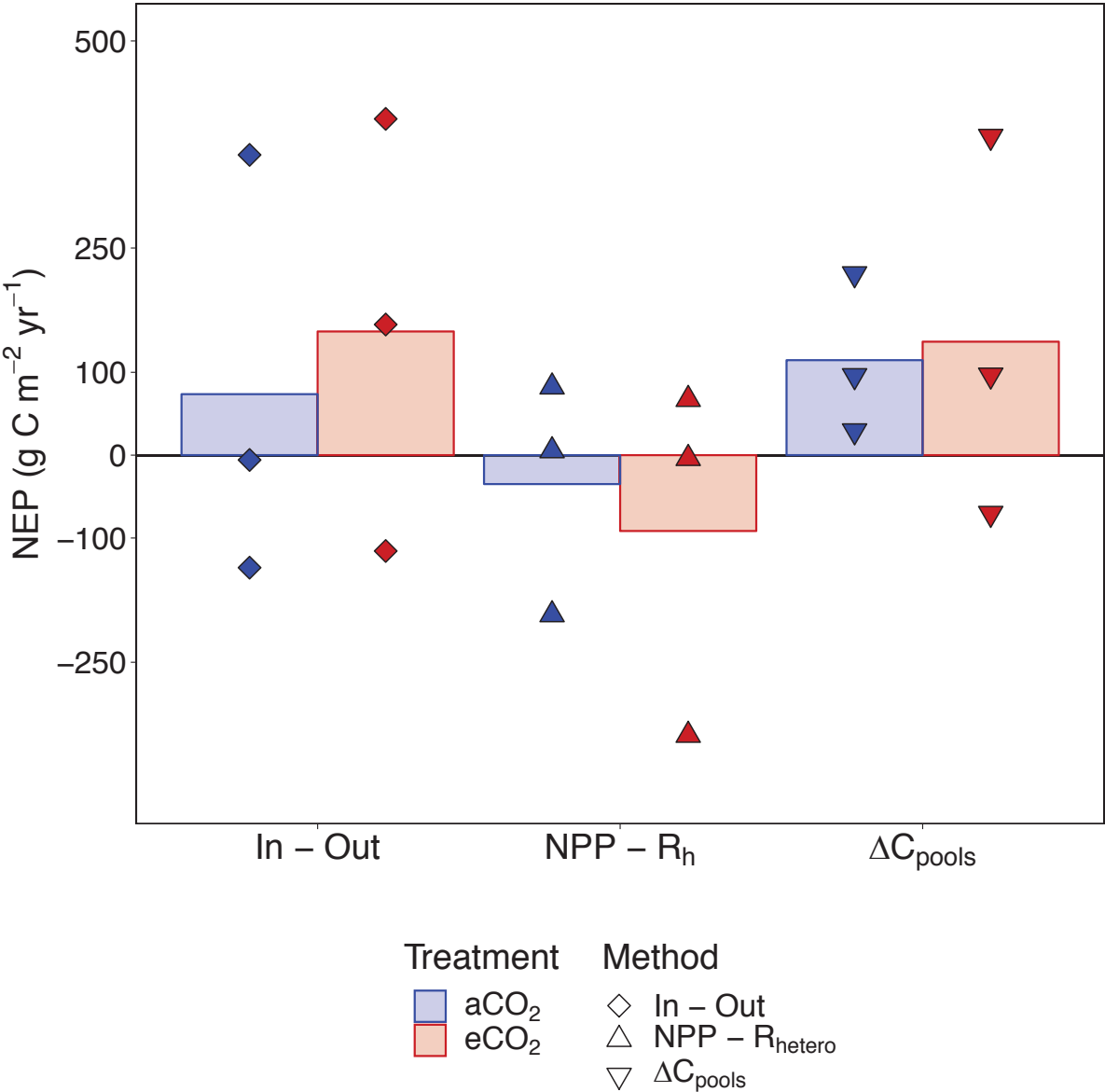
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782

783 **Figure 2. The fate of additional carbon fixed under elevated CO₂ (eCO₂) in a mature forest ecosystem. a)** Column “GPP” represents the total
784 eCO₂-induced increases in overstorey and understorey gross primary production (GPP_o and GPP_u, respectively), “NPP + R_a” represents the sum
785 of net primary production and autotrophic respiration response, “R + ΔC_{pools}” represents the sum of ecosystem respiration and carbon storage
786 response. **b)** The relative contributions of individual NPP fluxes to the aggregated NPP response to eCO₂, including NPP responses of overstorey
787 leaf (NPP_{ol}), twigs, barks and seeds (NPP_{other}), fineroot (NPP_{froot}), and understorey aboveground (NPP_{ua}); **c)** The relative contributions of individual
788 respiratory fluxes to the aggregated R response to eCO₂, including respiration responses of stem (R_{stem}), root (R_{root}), understorey aboveground

789 (R_{ua}), and soil heterotroph (R_{hetero}); and **d**) The relative contributions of individual change in carbon storage to the aggregated ΔC_{pools} response to
790 eCO_2 , including changes in pool of overstorey leaf (ΔC_{ol}), stem (ΔC_{stem}), understorey aboveground (ΔC_{ua}), fineroot (ΔC_{froot}), and soil (ΔC_{soil}).
791 Variables with an absolute mean CO_2 effect of $< 5 \text{ gCm}^{-2}\text{yr}^{-1}$ are excluded from the figure for better visual clarification. Individual CO_2 responses
792 are reported in Extended Data Figure 4. Each color represents the CO_2 response of a flux variable, point indicates the net sum of all variables for
793 a column, and the grey error bar represents one standard deviation of the estimated column sum at the plot-level (see Methods). The CO_2 effect is
794 estimated using a linear mixed-model analysis with plot-specific pre-treatment leaf area index as a covariate to account for pre-existing differences
795 (see Methods). The un-normalized response is provided in Extended Data Figure 3, which generally agrees with findings present in this figure, but
796 with less statistical precision.



798

799 **Figure 3. Estimates of net ecosystem production (NEP) under ambient and elevated CO₂**
800 **treatment at EucFACE.** Positive values indicate ecosystem net carbon uptake by the
801 ecosystem. “In - Out” calculates NEP based on the difference between total influxes and total
802 outfluxes. “NPP - R_{hetero}” calculates NEP based on the difference between net primary
803 production (NPP) and heterotrophic respiration (R_{hetero}). “ΔC_{pools}” derives NEP based on
804 incremental changes in all ecosystem carbon pools. Colored bars indicate treatment means
805 based on each method (n=3), with blue representing ambient and red representing elevated CO₂

806 treatment. Individual dots are plot-level NEP, derived based on different methods (see
807 Methods). Values are normalized by a linear mixed-model with plot-specific pre-treatment leaf
808 area index as a covariate to account for pre-existing differences. Horizontal dotted line indicates
809 NEP equals zero.

810 **Extended Data Table 1. Definition and data availability of variables.** Data availability
811 includes start and end year of data included in this study. Time points indicate the number of
812 data collections over the available data period. Within plot sub-replicate indicate the number
813 of replicates within each treatment plot. The detailed methods for estimating each variable is
814 provided in the Method section.

Variable		Data coverage			
Name	Symbol	Start year	End year	Time points	Within plot sub-replicate (plot ⁻¹)
Specific Leaf Area	SLA	2013	2016	50	3
Leaf Area Index	LAI	2012	2016	303	1
Soil bulk density	BK	2017	2017	2	3
Diameter at breast height	DBH	2013	2016	4	Individual tree
Overstorey leaf pool	C _{ol}	2012	2016	303	1
Understorey aboveground pool	C _{ua}	2015	2016	16	4
Overstorey stem C pool	C _{stem}	2013	2016	4	Individual tree
Fine root C pool	C _{froot}	2014	2016	6	4
Coarse root C pool	C _{croot}	2013	2016	4	Individual tree
Forest floor leaf litter C pool	C _{lit}	2013	2016	46	-
Microbial C pool	C _{micr}	2012	2015	15	4
Soil C pool	C _{soil}	2012	2014	11	4

Mycorrhizal C pool	C_{myco}	2015	2015	3	-
Insect C pool (aerial)	C_{ins}	2013	2016	43	8
Insect C pool (ground dwelling)	C_{ins}	2013	2015	5	4
Overstorey gross primary production	GPP_o	2013	2016	Annual	1
Understorey gross primary production	GPP_u	2013	2016	Annual	1
Overstorey leaf respiration	R_{ol}	2013	2016	Annual	1
Understorey leaf respiration	R_{ua}	2013	2016	Annual	1
Stem respiration	R_{stem}	2012	2016	Daily	3
Root respiration	R_{root}	2012	2015	Daily	-
Methane net flux	CH_4	2013	2016	35	7
Volatile C emission flux	VC	2013	2016	Daily	1
Insect herbivore respiration	R_{ins}	2012	2014	22	-
Dissolved organic C loss flux	DOC	2012	2014	12	4
Soil respiration	R_{soil}	2012	2015	Daily	8
Growth respiration	R_{grow}	2012	2016	Annual	1
Overstorey leaf net primary production	NPP_{ol}	2012	2016	49	8
Stem net primary production	NPP_{stem}	2012	2016	4	Individual tree

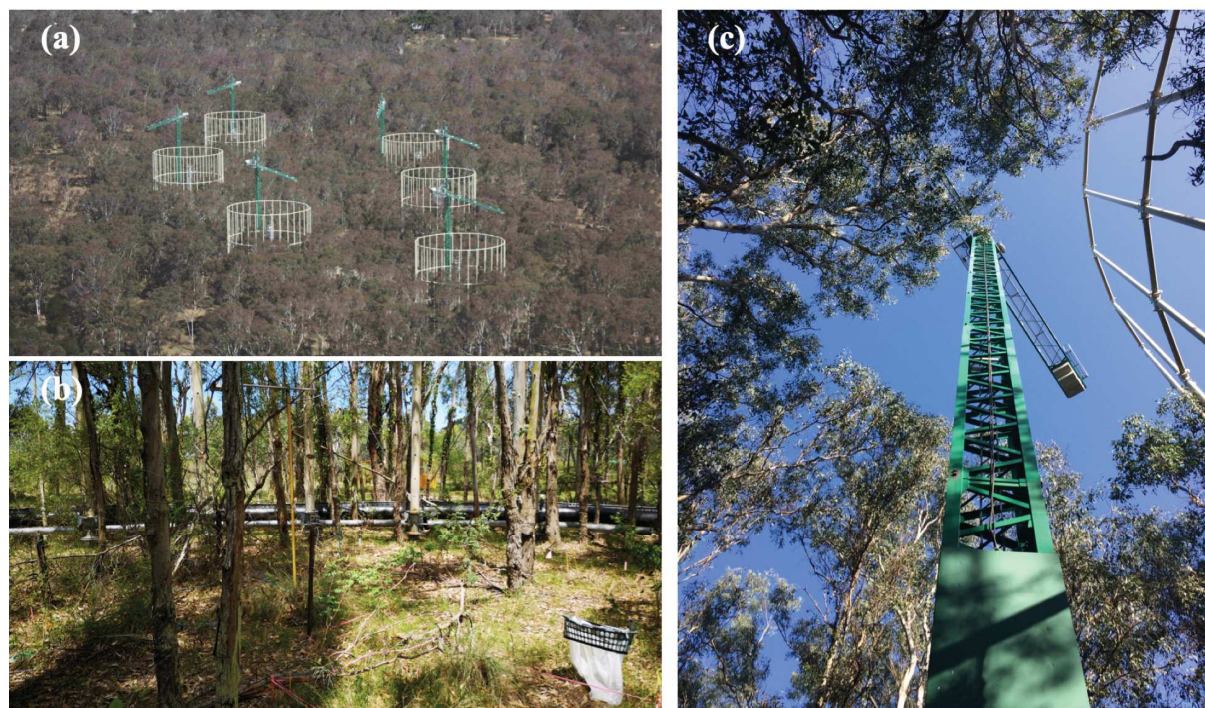
Fine root net primary production	NPP_{root}	2014	2016	5	4
Coarse root net primary production	NPP_{root}	2012	2016	4	Individual tree
Other net primary production (sum of twigs, bark, seeds)	NPP_{other}	2012	2016	49	8
Twig net primary production	NPP_{twig}	2012	2016	49	8
Bark net primary production	NPP_{bark}	2012	2016	49	8
Seed net primary production	NPP_{seed}	2012	2016	49	8
Understorey aboveground net primary production	NPP_{ua}	2015	2016	3	4
Frass production	Frass	2012	2014	22	8
Heterotrophic respiration	R_{hetero}	2012	2016	Daily	8
Overstorey leaf insect consumption flux	NPP_{ins}	2012	2014	22	-

816 **Extended Data Table 2. Carbon (C) fraction used to convert from biomass into C content.**

Variable	Symbol	Mean value	Data source
C fraction of overstorey leaf pool	f_{ol}	0.5	EucFACE data
C fraction of understorey aboveground pool	f_{ua}	0.456	EucFACE data
C fraction of stem pool	f_{stem}	0.445 (ambient plots) 0.448 (elevated plots)	EucFACE data
C fraction of coarse root pool	f_{croot}	0.445 (ambient plots) 0.448 (elevated plots)	Assumed the same as f_{stem}
C fraction of fine root pool	f_{froot}	0.40 (ambient plots) 0.42 (elevated plots)	EucFACE data
C fraction of overstorey leaf litter pool	f_{lit}	0.5	EucFACE data
C fraction of aboveground insect pool	f_{ins}	0.5	Ref 48
C fraction of frass production	f_{frass}	0.53	EucFACE data
C fraction of microbial pool	f_{micr}	0.534 (ambient plots) 0.493 (elevated plots)	EucFACE data

C fraction of mycorrhizal pool	f_{myco}	0.534 (ambient plots) 0.493 (elevated plots)	Assumed the same as f_{micr}
C fraction of soil pool	f_{soil}	0.016 (ambient plots) 0.017 (elevated plots)	EucFACE data
C fraction of twigs, barks and seeds production	f_{other}	0.5	Assumed

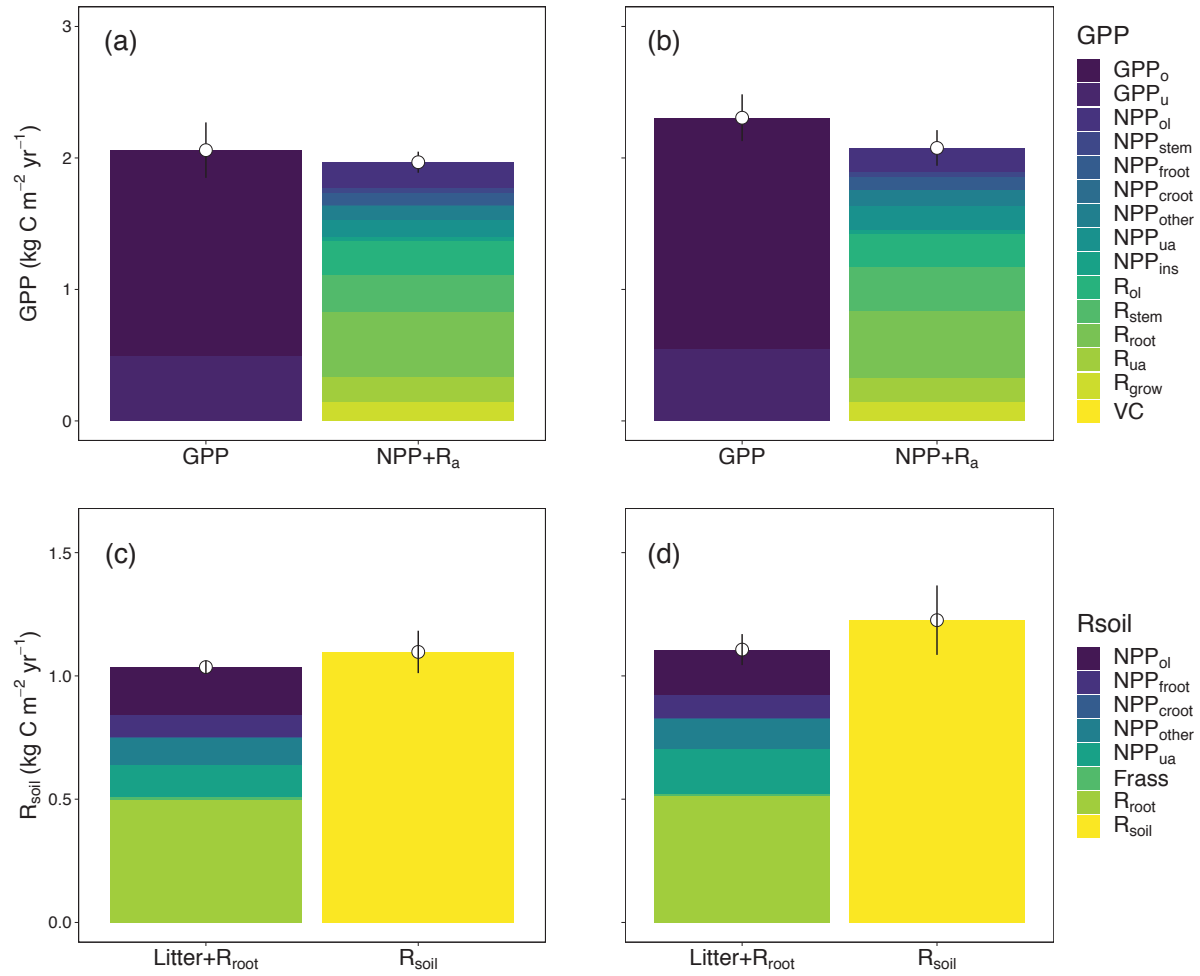
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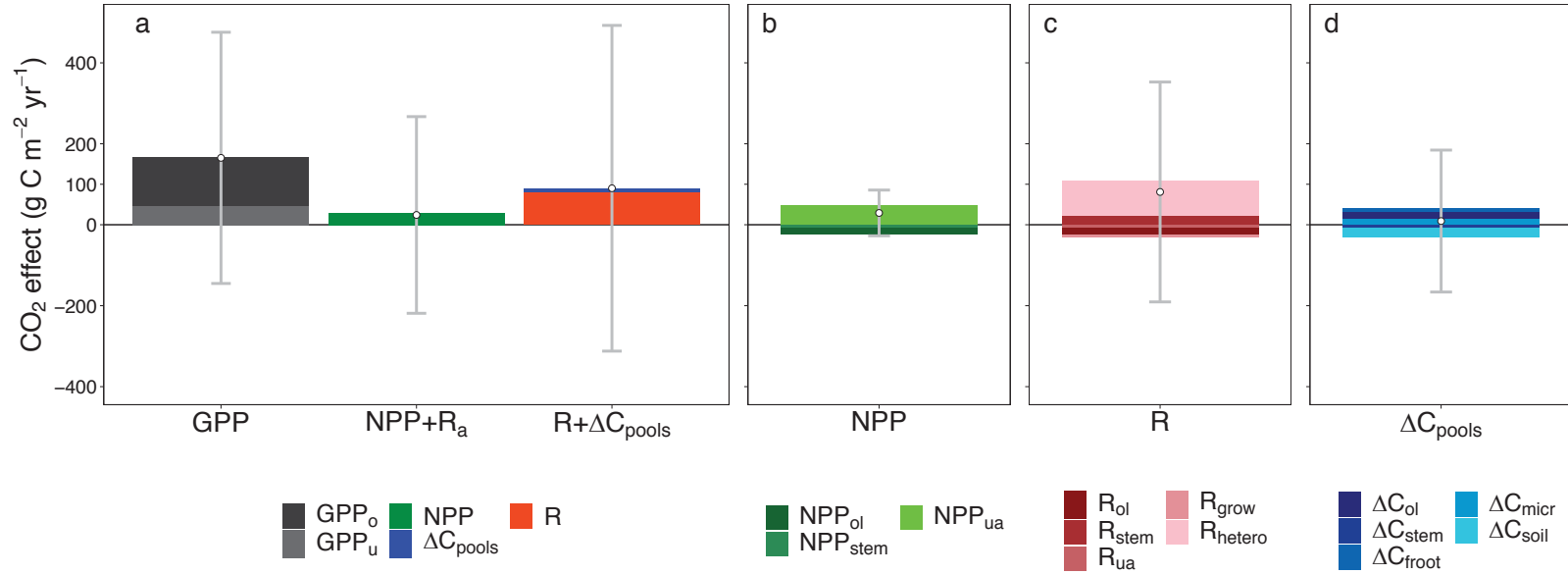
819 **Extended Data Figure 1. The *Eucalyptus* free air carbon dioxide enrichment experiment**
820 **facility (EucFACE). a)** A spatial overview of the forest and the facility (photo credit: David
821 S. Ellsworth), **b)** an overview of the understorey vegetation and infrastructure inside a plot
822 (photo credit: Mingkai Jiang), and **c)** a bottom-up look of the canopy structure and the crane
823 (photo credit: Mingkai Jiang).

824



827 **Extended Data Figure 2. Estimates of (a and b) gross primary production (GPP) and (c**
828 **and d) soil respiration (R_{soil}) based on different methods for both (a and c) ambient and**
829 **(b and d) elevated CO_2 treatment at EucFACE.** For estimates of GPP, we compared the
830 model simulated total GPP of overstorey and understorey (GPP_o and GPP_u , respectively), with
831 the sum of data-driven estimates of net primary production (NPP) and autotrophic respiration
832 (R_a), which include NPP of overstorey leaf (NPP_{ol}), stem (NPP_{stem}), fineroot (NPP_{froot}), coarse
833 root (NPP_{croot}), twigs, barks and seeds (NPP_{other}), understorey aboveground (NPP_{ua}), leaf
834 consumption by insects (NPP_{ins}), and respiratory fluxes of overstorey leaf (R_{ol}), stem (R_{stem}),
835 root (R_{root}), understorey aboveground (R_{ua}), growth (R_{grow}), and volatile carbon emission (VC).
836 For estimates of R_{soil} , we compared direct estimates of R_{soil} scaled up from soil chamber

837 measurements, with the sum of litterfall and independent estimates of root respiration ($\text{Litter} +$
838 R_{root}), assuming no net change in soil carbon stock over time. Here litterfall was inferred based
839 on NPP of overstorey leaf (NPP_{ol}), fineroot ($\text{NPP}_{\text{froot}}$), coarse root ($\text{NPP}_{\text{croot}}$), twigs, barks and
840 seeds ($\text{NPP}_{\text{other}}$), understorey aboveground (NPP_{ua}), and frass production (Frass). These
841 evaluations provide independent mass balance checks of the estimated ecosystem carbon
842 budget. Each color represents a flux variable. Dotted point and vertical line represent treatment
843 mean and standard deviation based on plot-level estimates of the aggregated flux ($n=3$). Values
844 were normalized by a linear mixed-model with pre-treatment leaf area index as a covariate to
845 account for pre-existing differences.



846

847 **Extended Data Figure 3. The fate of additional carbon fixed under elevated CO₂ (eCO₂) in a mature forest ecosystem (non-normalized**

848 **analysis case). a)** Column “GPP” represents the total eCO₂ induced increase in overstorey and understorey gross primary production (GPP_o and

849 GPP_u, respectively), column “NPP + R_a” represents the sum of net primary production and autotrophic respiration eCO₂ response, and column “R

850 + ΔC_{pools}” represents the sum of ecosystem respiration and carbon storage eCO₂ response. **b)** The relative contributions of individual NPP fluxes

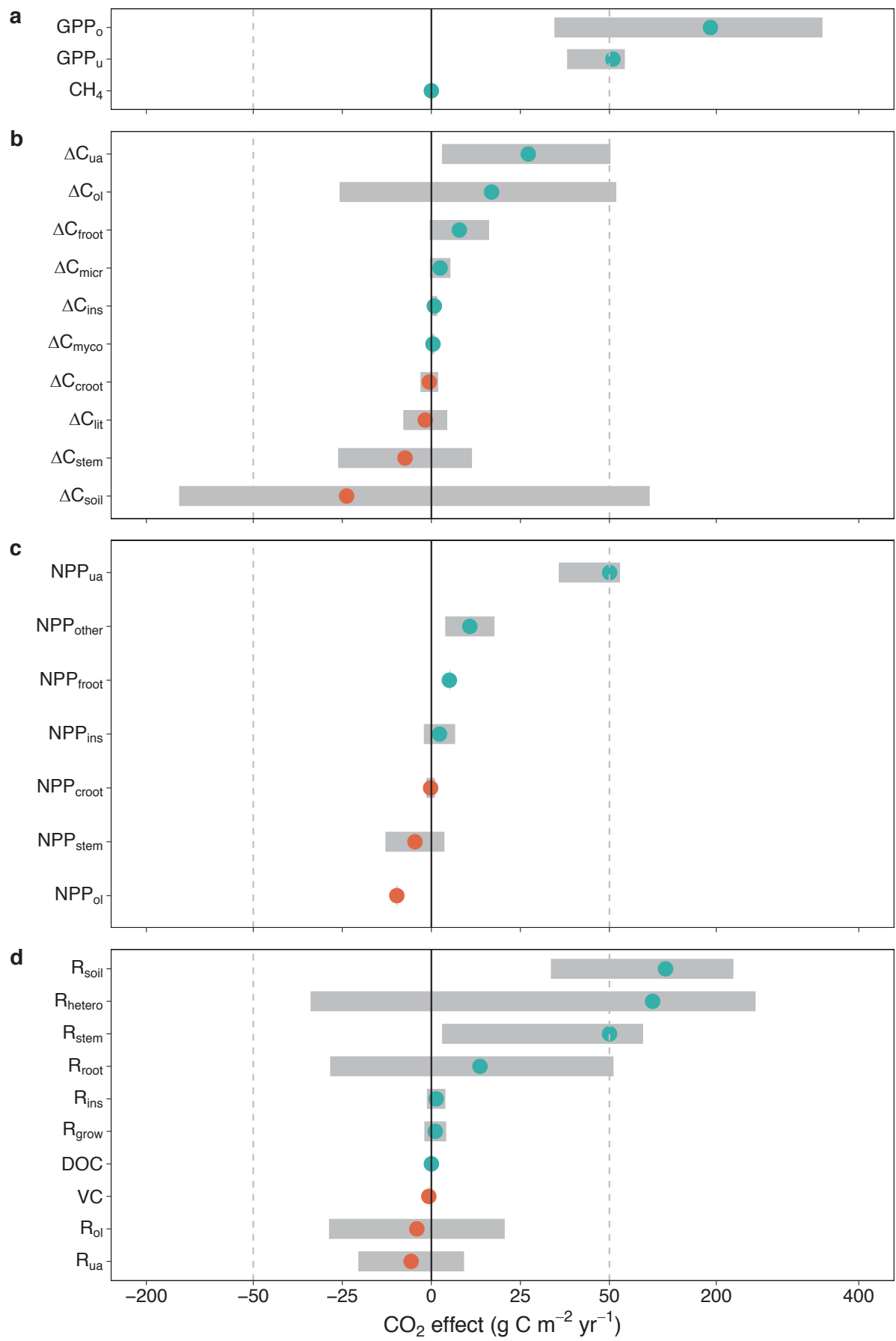
851 to the aggregated NPP response to eCO₂, including overstorey leaf (NPP_{ol}), stem (NPP_{stem}), and understorey aboveground (NPP_{ua}).

852 **c)** The relative contributions of individual respiratory fluxes to the aggregated R response to eCO₂, including overstorey leaf (R_{ol}), stem (R_{stem}), understorey

853 aboveground (R_{ua}), growth (R_{grow}), and heterotroph (R_{hetero}).

854 **d)** The relative contributions of individual change in carbon storage to the aggregated ΔC_{pools} response to eCO₂, including overstorey leaf (ΔC_{ol}), stem (ΔC_{stem}), fineroot (ΔC_{froot}), microbe (ΔC_{micr}), and soil (ΔC_{soil}). Variables with an

855 average CO₂ effect of $< 5 \text{ gCm}^{-2}\text{yr}^{-1}$ were excluded from the figure for better visual clarification. Each color represents a flux variable, point
856 indicates the net sum of all variables for a column, and the grey confidence interval represents plot-level standard deviation (n=3) of the estimated
857 column sum.



859 **Extended Data Figure 4. CO₂ treatment effect (gCm⁻²yr⁻¹) for all ecosystem fluxes at**
860 **EucFACE. a)** The CO₂ response of gross ecosystem carbon uptake, including gross primary
861 production of overstorey (GPP_o) and understorey (GPP_u), and soil methane uptake (CH₄). **b)**
862 The eCO₂ response of annual incremental change in carbon pool (ΔC_{pools}), including overstorey
863 leaf (ΔC_{ol}), stem (ΔC_{stem}), coarse root (ΔC_{croot}), fineroot (ΔC_{froot}), understorey aboveground
864 (ΔC_{ua}), leaf litter (ΔC_{lit}), soil (ΔC_{soil}), microbe (ΔC_{micr}), aboveground insect (ΔC_{ins}), and
865 mycorrhizae (ΔC_{myco}). **c)** The eCO₂ response of net primary production (NPP), including
866 overstorey leaf (NPP_{ol}), stem (NPP_{stem}), coarse root (NPP_{croot}), fineroot (NPP_{froot}), understorey
867 aboveground (NPP_{ua}), twigs, barks and seeds (NPP_{other}), and leaf insect consumption (NPP_{ins}).
868 **d)** The eCO₂ response of ecosystem respiration (R) and other out-going flux, including
869 respiration fluxes of overstorey leaf (R_{ol}), stem (R_{stem}), root (R_{root}), understorey aboveground
870 (R_{ua}), growth (R_{grow}), insect (R_{ins}), heterotroph (R_{hetero}), and soil (R_{soil}), and volatile carbon
871 emission (VC) and dissolved organic carbon leaching (DOC). Dots and grey bars represent
872 means and standard deviations of the CO₂ treatment difference, predicted by a linear mixed-
873 model with plot-specific pre-treatment leaf area index as a covariate. Orange dots indicate
874 negative means and light green dots indicate positive means. Dashed lines indicate change of
875 scale along the x-axis.

876