- Letter -

Recovery of trees from drought depends on belowground sink control

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1	Climate projections predict higher precipitation variability with more frequent dry extremes ¹ . CO ₂ -
2	assimilation of forests decreases during drought, either by stomatal closure ² or by direct
3	environmental control of sink tissue activities ³ . Ultimately, drought effects on forests depend on
4	the ability of forests to recover, but the mechanisms controlling ecosystem resilience are
5	uncertain ⁴ . Here, we investigated the effects of drought and drought release on the C balances in
6	beech trees by combining CO_2 -flux measurements, metabolomics and $^{13}CO_2$ -pulse labelling. During
7	drought, net-photosynthesis (A_N), soil respiration (R_s) and the allocation of recent assimilates
8	below ground were reduced. Carbohydrates accumulated in metabolically resting roots but not in
9	leaves, indicating sink control of the tree C balance. After drought release, R_s recovered faster than
10	A_N and CO ₂ -fluxes exceeded those in continuously watered trees for months. This stimulation was
11	related to greater assimilate allocation to and metabolization in the rhizosphere. These findings
12	show that trees prioritize the investment of assimilates below ground, probably to regain root
13	functions after drought. We propose that root restoration plays a key role in ecosystem resilience
14	to drought, in that the increased sink activity controls the recovery of C balances.
15	Forests play a crucial role in the global carbon (C) cycle because they hold a large fraction of the
16	global C stock and act as a major sink for atmospheric $\rm CO_2^5$. However, drought reduces primary
17	productivity, thereby turning forests from C sinks into C sources ⁶ . It has generally been assumed that
18	plant and ecosystem C balances under drought are controlled by restricted photosynthetic source
19	activity rather than by changes in the sink activity of plant tissues ^{7,8} . Recently, direct environmental
20	control of sink activity with feedbacks to CO_2 -assimilation has been proposed ⁹ , but no unequivocal
21	evidence has been obtained yet. Of comparable importance, but even less understood, are the
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	mechanisms controlling plant and ecosystem C balances after drought release, though the ability of
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23 24 25	mechanisms controlling plant and ecosystem C balances after drought release, though the ability of plants to restore CO ₂ -assimilation and other functions determines the resilience of trees and forest ecosystems. Further, limited knowledge on the principles that control C allocation in trees prevents us from predicting C balances of forests under future environmental conditions characterized by

27	Using two experimental setups, with beech growing either in model ecosystems in open-top
28	chambers (Supplementary Figure 1) or in pots, we studied tree and ecosystem C fluxes during
29	drought and after drought release. By combining measurements of net-photosynthesis (A_N) and soil
30	respiration (R_s) as indicators of source and sink activity, respectively, with ¹³ CO ₂ -pulse labelling and
31	metabolomic analyses, we followed seasonal C dynamics and tracked assimilate transport through
32	the plant-soil system. Based on a hypothetical framework (Figure 1), we aimed to test if changes in
33	A_N and R_s , as well as shifts in carbohydrate allocation, indicate source or sink control of C balances. If
34	source activity controls C balances under drought, we expected an initial decrease in A_N and leaf
35	carbohydrate concentrations and a delayed depletion of carbohydrates in roots, leading to a
36	reduction in R_s (Figure 1a). A comparable response would occur upon drought release, with an initial
37	recovery of A_N and leaf carbohydrate concentrations followed by a delayed increase of R_s (Figure 1b).
38	If, however, the C balance is sink-controlled, drought would directly reduce R_s , leading to an
39	accumulation of carbohydrates in roots due to reduced C demand. In this case, A _N would acclimate to
40	the reduced sink demand after a delay and leaf carbohydrate concentrations consequently would not
41	change (Figure 1c). Upon drought release, R_s would increase and, with a delay, the increased
42	belowground C demand would positively feed back on A_N (Figure 1d).
43	In our model ecosystems, A_N and R_s decreased by 44% and 28%, respectively, over the entire drought
44	season (Figure 2a,b,c; Supplementary Figure 2a). At the end of the drought, a ¹³ CO ₂ -pulse label was
45	applied to the canopies to trace the fate of recent assimilates in the plant-soil system. Under
46	drought, the uptake of 13 CO $_2$ decreased by 81% and assimilate translocation to belowground sinks
47	was reduced, as shown by lower ¹³ C signals in mycorrhizal roots and soil microbial biomass and
48	reduced 13 CO ₂ soil efflux (Figure 3a,c,e,g). The reduction in 13 CO ₂ soil efflux was 83% and thus
49	comparable to that in 13 CO ₂ uptake. However, the 13 CO ₂ soil efflux showed a stronger reduction (83%)
50	than that observed for R_s (approx. 50%), indicating that other C sources, either related to
51	heterotrophic soil respiration or tree internal C storages ¹⁰ , contributed to soil respiration but were
52	less sensitive to drought. Under drought, the 13 C peak in continuously monitored soil CO $_2$

53 (Supplementary Figure 3a) was delayed by one day and mean residence times (MRT) of assimilated 54 ¹³C in the plant-soil system increased (drought 76 h, control 30 h), indicating slower assimilate transport to belowground sinks, as previously reported¹¹. The reduced and delayed assimilate 55 56 transport might have been the result of either source limitation or sink control. However, non-57 structural carbohydrates (NSC) were not depleted in source leaves in drought-treated model 58 ecosystems (Supplementary Table 1) and thus source limitation was unlikely, as recently proposed⁹. 59 To explore the mechanisms leading to reduced C fluxes to belowground sinks, we studied the 60 dynamics of metabolites with progressing drought in a pot experiment. Reductions in soil moisture 61 and A_N in drought-treated pots were comparable to those in the model ecosystems (Figure 4a,b; 62 Supplementary Figure 2b). In roots, the NSCs fructose, glucose, sucrose and starch, as well as the 63 osmoprotectant proline, increased under drought and sucrose accumulated in the release phloem. 64 The increase in the concentration of NSCs by up to 700% was very strong but still in the range reported in previous studies with trees^{12,13}. In leaves, no NSC increase was observed except for a 65 66 delayed accumulation of starch and proline as drought progressed. Although NSCs accumulated in 67 roots, sink control of such an increase can only be inferred when the size of carbohydrate pools 68 depends directly on the balance between supply through photosynthesis and demand for growth and respiration¹⁴. Alternatively, accumulation of NSC might serve as osmotic adjustment¹⁵, which is not 69 70 directly related to changes in C supply and demand. In our study, however, accumulation of NSC was only observed in roots, whilst proline, an indicator of osmotic regulation¹⁶, was enriched in both 71 72 roots and leaves. It is thus unlikely that osmotic adjustment via an increase in NSC content was 73 achieved only in roots and not in leaves. Instead, active storage of carbohydrates in roots at the 74 expense of metabolic processes might have occurred, as previously suggested¹⁷. Thus, the strong 75 accumulation of NSC in roots and the lack thereof in leaves reflect metabolic activity in sink and 76 source tissues. Transferring this information to our model ecosystems indicates that the reduced C 77 flux to belowground sinks under drought (Figure 3c,e,g) was a consequence of decreased sink activity 78 (Figure 1c). Due to the rather slow build-up of drought over time in our model ecosystems, a clear

order of the response of source (A_N) vs. sink (R_s) activities could not be derived directly, especially since changes in R_s might have been only partially due to changes in autotrophic root-rhizosphere respiration.

82	The recovery of plant and ecosystem C fluxes after prolonged drought was examined by re-watering
83	the model ecosystems. R _s responded rapidly, reaching control values within the first three days and
84	exceeding thereafter values in controls until the end of the growing season (Figure 2b,c;
85	Supplementary Table 3). This stimulation of R_s nearly compensated for the previous drought
86	reduction, with the flux integrated over the entire growing season amounting to 98% of that in
87	controls. A_N responded similarly but the recovery was delayed by approx. one week due to metabolic
88	limitation, as shown by impaired PSII photochemistry (Figure 2a,c; Supplementary Table 3). Further,
89	the stimulation of A_N occurred later and compensated for only 82% of the previous drought
90	reduction. A second 13 CO ₂ -pulse label was applied to the tree canopies when R_s exceeded the values
91	in controls but A_N was still slightly below that of controls. The previous drought exposure increased
92	the translocation of recent assimilates to belowground sinks compared to controls, as shown by
93	higher 13 C signals in mycorrhizal roots and soil microbial biomass and by an enhanced 13 CO $_2$ soil efflux
94	(Figure 3b,d,f,h). The latter signal was enhanced by 50% and thus increased in relation to
95	photosynthetic ¹³ C uptake, which was not affected. This increase represents exclusively autotrophic
96	respiration, and the comparable increase in R_s (68% R_s vs. 50% ¹³ CO ₂ soil efflux) shows that the plant-
97	driven C flux was primarily responsible for the observed stimulation of R_s . Heterotrophic soil
98	respiration was small in our model ecosystems containing low soil organic content and can therefore
99	be excluded as the cause of stimulated R_s because it only responds transiently to rewetting of dry
100	soils by the so-called "Birch Effect" ^{18,19} . Our results thus clearly show that a drought effect is
101	imprinted on plant source and sink tissues, supporting the concept of an ecological stress memory of
102	which the underlying mechanisms are still poorly understood ²⁰ .
103	The increased C demand of belowground sinks resulted in only a slight feedback on the velocity of C

104 transport. While the peak time of ${}^{13}CO_2$ soil efflux was comparable, the *MRT* of ${}^{13}C$ in the plant-soil

105 system was somewhat lower for the previously drought-exposed trees, as calculated from 106 continuously monitored soil CO₂ (post-drought 36 h, control 41 h; Supplementary Figure 3b). 107 However, the mass flow of assimilates to belowground sinks can additionally be increased if less C is 108 unloaded from the transport pathway for storage or growth in aboveground tree organs²¹. Indeed, 109 the C allocation to growth in twigs, stem and roots did not fully recover after re-watering, indicating 110 that growth-related sink activity along the transport pathway was still reduced (Supplementary Table 111 4). The greater allocation and use of recent assimilates in belowground sinks after re-watering shows 112 that trees give high priority to investing into their roots upon recovery from drought. The likely 113 reason for this response is the metabolic need for root and mycorrhizal restoration in order to restore trees' capability to acquire water and nutrients after an extended drought^{22,23}. Effects on root 114 115 growth can be excluded, as demographic root characteristics were not affected during or after 116 drought (Supplementary Figure 4a-c). Thus, root and mycorrhizal restoration relied mainly on 117 increased metabolic activity, which explains the fast recovery and stimulation of R_s. Since A_N showed 118 a delayed recovery and later stimulation than R_s, the latter was clearly not source driven and instead 119 reflects the metabolic need for root and mycorrhizal restoration. On the contrary, we postulate that 120 increased belowground sink activity upon drought release feeds back on A_N , triggering the delayed 121 recovery and stimulation of CO₂-assimilation (cf. Figure 1d). Furthermore, our findings support sink 122 control of the C balance under previous drought conditions, as a drought-induced depletion of 123 belowground C reserves should delay the recovery of R_s compared to A_N if the recovery is source 124 controlled.

There is increasing evidence that drought not only influences ecosystem C balances concurrently but also triggers delayed responses that involve multiple mechanisms operating at different scales of time, plant function and ecosystem organization^{4,24}. To date, such mechanisms are poorly understood and thus constitute a large uncertainty in projections of ecosystem C balances and resilience. Here, we show that tree C fluxes not only recover but even increase after drought to compensate for previous stress impacts. This compensation is sink-driven, leading to a greater

131	belowground allocation of recent assimilates upon drought release. The observed response has
132	important consequences for ecosystem C cycling, as it increases the input of plant-derived labile C
133	into soils, thereby fueling soil microbial communities ²⁵ . We suggest that the ability of trees to
134	reactivate root metabolism is vital for ecosystem resilience to drought. However, the extent of this
135	effect very likely depends on the severity and duration of drought and may vary with tree age, as
136	adult trees have larger C storage compartments. Taken together, our findings suggest a resilience
137	mechanism that attenuates drought disturbances of seasonal tree C balances and needs to be
138	considered when estimating the impact of climate change on the C balances of forest ecosystems.

139 MATERIAL AND METHODS

140 Plant material and growth conditions

- 141 The model ecosystem experiment was conducted in 16 field-based open top chambers. In each
- 142 chamber, a model ecosystem was established with young beech trees (Fagus sylvatica L.) growing on
- 143 lysimeters filled with forest soil of low soil organic C content (Supplementary Figure 1;
- 144 Supplementary Methods; Supplementary Table 2). A summer drought was simulated by reducing the
- 145 water supply from 22 May to 1 August by 78%. After the trees had developed the critical water
- 146 deficit for leaf physiological function (predawn water potentials below -2 MPa^{27,28}), the lysimeters
- 147 were intensely re-watered and afterwards regularly irrigated until the end of the vegetation season
- 148 (Supplementary Figure 2a).
- 149 The pot experiment was carried out with beech saplings (*Fagus sylvatica* L.) in a greenhouse
- 150 environment (Supplementary Methods). During the drought treatment lasting 4 weeks, the control
- 151 pots were watered to field capacity while pots with the drought treatment received no water at all.
- 152 Fine root and leaf samples were taken weekly.

153 Measurements of net-photosynthesis and soil respiration

- 154 Net-photosynthesis (A_N) was measured on 3-4 trees per lysimeter between 11:00 and 16:00 CET
- using a photosynthesis system (LI-COR 6400, Lincoln, USA) equipped with a broadleaf cuvette. The
- 156 conditions inside the cuvette were kept constant at 400 ppm CO_2 and a photon flux of 1000 μ mol m⁻²
- 157 s⁻¹. Metabolic constraints on A_N were tested by chlorophyll fluorescence analysis using the
- 158 performance index PI_{total} of $PSII^{29}$. Soil respiration (R_s) was measured with a custom-made static
- 159 chamber³⁰ equipped with a diffusion-aspirated non-dispersive infrared analyser connected to a
- 160 humidity/temperature sensor (GMP343 CO₂ probe, HMP75 rH/T probe; Vaisala, Vantaa, Finland). The
- 161 increase in CO₂-concentrations in the chambers was measured in permanently installed PVC collars (5
- 162 cm height, two per lysimeter).

163 ¹³C pulse labelling

164 Allocation of assimilates was followed by ¹³C-pulse labelling in six randomly selected lysimeters (n = 3 165 per treatment) at the end of the drought and in eight lysimeters (n = 4 per treatment) two weeks after re-watering. Before labelling, the soil was covered with plastic foil to minimize diffusion of ¹³CO₂ 166 167 into the soil. All trees in a given lysimeter were covered with a tall tent made of transparent plastic 168 foil. The CO₂-concentration inside was reduced to 200 ppm by flushing the tent with CO₂-free air. The labelling lasted 2 h, during which time we added 100% CO₂ with a 50:50 ratio of 13 CO₂ and 12 CO₂. The 169 170 CO_2 -concentration was kept constant at about 1500 ppm, which is above the saturation point for 171 CO₂-uptake. 172 ¹³C analysis in leaves, mycorrhizal root tips, soil microbial biomass and soil-respired CO₂ 173 Leaves from 3-4 trees per lysimeter were oven-dried at 60°C, milled and weighed into tin capsules for 174 ¹³C analyses. Mycorrhizal root tips and soil microbial biomass were randomly sampled in each 175 lysimeter in the upper 10 cm soil depth by taking three soil cores with a diameter of 2 cm. Additional 176 roots were taken directly from 3-4 trees. Vital mycorrhizal root tips were immediately collected 177 under a stereomicroscope and kept at -70°C until processing. They were pooled per lysimeter, ovendried at 80°C, milled and weighed into tin capsules for ¹³C analyses. Soil microbial biomass was 178 179 determined using the chloroform fumigation extraction method, whereby the concentration and 180 isotopic signature of extracted organic C from non-fumigated and fumigated samples were determined by oxidizing extractable C to CO_2^{31} . The ¹³C of microbial biomass was calculated as 181 described previously³². The ¹³C signature of soil-respired CO₂ was determined by the closed chamber 182 method³³. For each sample, the collars were closed with 7 cm tall PVC lids with cellular rubber and 183

184 gas samples were taken after 15 min. In addition, ambient air close to the soil surface was collected185 at each sampling occasion.

186 In gas samples, the δ^{13} C values and the CO₂-concentration were analysed with a GasBench II coupled 187 to a Delta V Plus mass spectrometer (ThermoFinnigan, Bremen, Germany). The ¹³C signatures in solid 188 samples were measured with an Elemental Analyzer (Euro EA, Eurovector, Milano, Italy) coupled to

the mass spectrometer. The δ^{13} C value of soil-respired CO₂ was calculated as a mixture of ambient and soil-respired CO₂ sampled in the chamber³⁴. The ¹³C signal (Δ^{13} C) in mycorrhizal roots was the difference between δ^{13} C values during and before labelling. The amount of ¹³C assimilated by plants, in soil microbial biomass and in soil-respired CO₂ was estimated by first expressing the δ notations in atom% and then calculating the excess ¹³C values considering each pool and flux size¹¹ (Supplementary Methods). The mean residence time (*MRT*) for the ¹³C soil efflux was calculated as

195 described previously¹¹.

196 Analysis of metabolites

197 Metabolites were analysed according to previous studies^{35,36}. In brief, frozen tissue was homogenized

and extracted with 87% methanol. Phloem exudates were obtained as previously described³⁷, dried

and re-dissolved in 87% methanol. Aliquots were derivatized and injected into a GC-quadrupole MS

200 system (GC: 7890A; MS: 5975C; Agilent Technologies, Waldbronn, Germany). GC-MS data were then

201 deconvoluted, peak areas quantified and mass spectra identified according to³⁵. Relative

202 concentration changes were calculated as log₁₀ ratios between drought and control treatments.

203 Statistical analysis

204 Data were analysed by fitting linear mixed effects models using maximum likelihood (lme function;

205 nlme package, R version 3.1.2.)³⁸ (Supplementary Table 3). For the entire measurement period,

season (dry: 22 May to 1 Aug vs. wet: 2 Aug to 31 Oct), treatment (drought/post-drought vs. control)

207 and date of measurement were used as fixed effects and lysimeter and individual tree were included

as random effects. The corAR1 function was included in the model to account for repeated

209 measurements with a first-order autoregressive covariate structure. Treatment effects were

additionally analysed for dry and wet season. To account for the varying ¹³C signal in the

211 consecutively labelled lysimeters, we included a co-variate as a fixed effect, thereby normalizing the

212 ¹³C tree uptake in each lysimeter to the treatment mean of the wet and dry season, respectively,

213 which allowed us to consider the treatment-specific ¹³C uptake by trees. In all final models, normality

- and homoscedasticity of the residuals were verified with diagnostic plots and the dependent
- 215 variables were all log or square root transformed.

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312 AUTHOR CONTRIBUTIONS

- 313 M.A., R.S., F.H., M.S. and A.G. designed the experiments; R.S., F.H., J.J. and M.A. performed the ¹³C-
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- 316 F.L. and M.L. analysed metabolites; J.J. and F.H. performed statistical analysis; A.G., J.J., F.H. and
- 317 M.A. wrote the manuscript.

318 COMPETING INTERESTS

319 The authors declare no competing financial interests.

320 FIGURE LEGENDS

321 Figure 1: Hypothetical trajectories of metabolic activity and metabolite concentration in leaves and 322 roots as a consequence of drought onset and drought release. For both drought (a, c) and drought 323 release (b, d), the scenarios for full source (a, b) and full sink (c, d) control of the tree carbon balance 324 after the change in soil moisture conditions are shown. Effects in source (leaves/green) and sink 325 tissues (root/brown) are provided for each scenario. We refer to net-photosynthesis (A_N) as a source 326 metabolic activity and to soil respiration (R_s) as an integrator of sink metabolic activity in the roots. 327 Metabolite content refers to the most abundant carbohydrates (i.e. the non-structural carbohydrates 328 (NSC) glucose, fructose, sucrose and starch; see also Figure 4b). Under source control (a, b), source 329 metabolic activity in leaves (A_N) reacts first to changing conditions and induces changes in assimilate 330 (i.e. sugar) availability for sinks and thus affects sink metabolic activity in roots (R_s) . Under sink 331 control (c, d), sink metabolic activity is directly affected by the environmental conditions, leading to 332 changes in sink metabolite levels. After a delay, source metabolic activity is impacted in response to 333 the altered sink demand. 334 Figure 2: Reduction of net-photosynthesis (A_N) and soil respiration (R_s) in the model ecosystem 335 experiment during drought (dry season), and during recovery and stimulation after drought release 336 (wet season). a, and b, show effects on A_N and R_S , respectively. Numerical values provide 337 quantitative measures of the drought limitation and the stimulation after full recovery ($P < 0.05^*$, 338 0.01^{**} , 0.001^{***} ; means ± SE; A_N : n = 8, R_S : n = 3-8). In **c**, the development of drought and post-339 drought effects on A_N (green) and R_S (brown) are shown, together with the release of metabolic

- 340 limitation of A_N after re-watering (PS II chlorophyll fluorescence; n = 8). Responses are shown as
- 341 relative deviations from control values.

Figure 3: Suppressed uptake and allocation of ¹³C assimilates in the model ecosystem experiment 342 343 under drought (dry season) and increased transfer to and metabolization in the belowground compartment after drought release (wet season). a, and b, leaf photosynthetic uptake of ¹³C. c, and 344 d, incorporation of ¹³C into mycorrhizal root tips. e, and f, transfer of ¹³C to soil microbial biomass. g, 345 and **h**, respiratory ¹³C release from the soil including mean residence times of recent assimilates 346 347 calculated from the δ^{13} C of continuously measured soil CO₂ (Supplementary Figure 3). *P* values < 0.05 348 indicate statistically significant treatment effects (means \pm SE, n = 3 for dry season and n = 4 for wet 349 season).





Figure 1



Figure 2



Figure 3



Figure 4