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Climate warming and elevated CO₂ alter peatland soil carbon sources and stability

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Peatlands are an important carbon (C) reservoir storing one-third of global soil organic carbon (SOC), but little is known about the fate of these C stocks under climate change. Here, we examine the impact of warming and elevated atmospheric CO₂ concentration (eCO₂) on the molecular composition of SOC to infer SOC sources (microbe-, plant- and fire-derived) and stability in a boreal peatland. We show that while warming alone decreased plant- and microbe-derived SOC due to enhanced decomposition, warming combined with eCO₂ increased plant-derived SOC compounds. We further observed increasing root-derived inputs (suberin) and declining leaf/needle-derived inputs (cutin) into SOC under warming and eCO₂. The decline in SOC compounds with warming and gains from new root-derived C under eCO₂, suggest that warming and eCO₂ may shift peatland C budget towards pools with faster turnover. Together, our results indicate that climate change may increase inputs and enhance decomposition of SOC potentially destabilising C storage in peatlands.

The capacity of peatlands to store more than one-third of the global SOC pool under waterlogged conditions^{1,2} has been touted as a powerful natural form of C sequestration³. However, there is concern that rising global temperatures and atmospheric CO₂ concentrations, along with corresponding changes in hydrology and biology, have the potential to destabilize SOC stocks and increase the flux of CO2 and CH4 from peat soils to the atmosphere^{2,4}, amplifying the drivers of climate change. Yet processes regulating the formation and stability of peat SOC, and their responses to interactive climate change factors remain unclear^{5,6}, making it difficult to forecast changes in C dynamics in peatland ecosystems⁷. In peatlands, SOC stability is controlled by a suite of environmental conditions, including temperature, hydrology (water saturation), and pH^{8,9}, and biotic factors such as plant and microbial community composition and functioning^{10,11} as well as the chemical characteristics of the peat itself^{12,13}. These biotic and abiotic factors regulate the current rates and pathways of enzymatically catalysed SOC decomposition^{8,10}, and they are expected to determine the response of SOC decomposition under future environmental change^{5,7}.

A key question for future peatland C balance is whether a portion of SOC that is currently stable may become more accessible to microbial decomposition^{2,4,7}, via direct effects of rising temperature and CO₂ concentration, indirect effects of altered water table^{9,14,15}, or due to changes in plant production and thus litter inputs to SOC^{11,16,17}. Soil warming, occurring more rapidly at high compared to low latitudes^{18,19}, can lead to lower water-tables and increases in peat aeration^{9,15}. Changes in moisture availability combined with the fertilization effects of elevated CO₂ can alter plant biomass production and the quality of litter inputs^{10,16,20}. Such a change in peat aeration and plant litter (including root exudates) inputs can impact the microbially-mediated SOC turnover because soil water content and litter chemistry regulate microbial community composition, function, and enzymatic activity⁸⁻¹¹. Differences in litter chemistry can also affect

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the sensitivity of organic matter decomposition to soil temperature, with further effects on C cycling^{10,21}. Consequently, if altered organic matter sources persist over extended periods, they are likely to stimulate the mineralisation of currently stable C²². In short, climate driven changes in water-table and plant litter inputs are likely to modify C storage^{5,6}, but the underlying mechanisms and their interactions, particularly those resulting in shifts in organic matter inputs and composition are still unknown in peatlands⁷. The role of such plant-soil interactions in regulating future SOC storage merits attention because a much larger proportion of SOC may be susceptible to climate-mediated losses than previously assumed^{7,19}.

Here, we investigate the effects of warming and elevated atmospheric CO₂ concentration (eCO₂) on peatland SOC molecular composition to infer SOC sources, stability, and implications for storage. To describe SOC dynamics, we partitioned bulk soil C into classes of molecular compounds comprising SOC defined operationally by organic matter origins (plant-, microorganism- and fire-derived) and potential decomposition rates²³⁻²⁵. Specifically, we targeted SOC compounds with distinct turnover times. Firstly, we targeted solventextractable compounds (alkanoic acids, alkanols, alkanes, steroids, and terpenoids, derived from plant- and microbial-material²⁶) that are thought to turn over predominantly on a timescale of bulk SOC or faster^{24,27}. Secondly, we targeted SOC components from different plant origins, hydrolysable biopolymers distinct to either leaf/needle (cutin) or root/bark (suberin) compartments (and presumably slowly cycling)²⁶⁻²⁹. Lastly, we targeted lignin and pyrogenic carbon (PyC) representing the most slowly cycling $C^{23,30-32}$. We assessed the stability of these SOC compound classes following 4 years of warming and 2 years of eCO₂ at an in-situ climate manipulation experiment (SPRUCE: Spruce and Peatland Responses Under Changing Environments). The SPRUCE experiment provides a powerful climate change gradient whereby above- and below-ground warming (whole-ecosystem warming; +0, +2.25, +4.5, +6.75, +9 °C above ambient) are crossed with ambient and elevated CO₂ concentrations (+500 ppm above ambient)³³. Given that high-latitude soils are projected to experience temperature increases of up to 8.3 ± 1.9 °C by 2100 under a high CO₂ emission scenario (RCP8.5)19, this unique experimental set up allowed us to explore the fate of SOC across mild to extreme future warming scenarios. From the perspective of long-term peatland C sequestration, we are particularly interested in the response of polymeric SOC components because kinetic theory predicts that decomposition and turnover of complex SOC molecules (greater chemical stability) should be more sensitive to temperature changes than simple molecules³⁴. In peatlands, warming is typically accompanied by a lower water-table coupled with peat aeration, which can further impact the apparent temperature sensitivity of organic matter decomposition^{8,9,14,15}. Consequently, if temperature and oxygenation effects predominate over other biotic and abiotic factors that influence decomposition dynamics^{9,25,34}, we hypothesize that complex SOC molecules such as lignin, and PyC that have been proposed to turn over much slower than bulk C in soils due to their polyaromatic chemical structure^{23,30,35}, would degrade faster. Overall, our study rejects this hypothesis that complex SOC molecules are more responsive to temperature changes. Instead, we found that all molecular compounds comprising SOC, regardless of source and complexity, were vulnerable to shifts in climate drivers, demonstrating the high sensitivity of peatlands to climate change.

Results and discussion Warming and eCO₂ have divergent effects on plant- and microorganism-derived SOC

First, we assessed how whole-ecosystem warming under ambient CO_2 modified SOC molecular composition. Four years of experimental warming decreased SOC molecules content in the surface peat (0–30 cm; Fig. 1; Fig. S2). Solvent-extractable compounds of plant and

microbial origins^{26,27,29} declined by 30% between the 0 and 9 °C treatments which corresponds to a slope of $-0.79 \pm 0.2 \text{ mg g}^{-1}$ per °C increase in temperature ($r^2 = 0.60$, p = 0.0004; Fig. 1a, Fig. S2a). Similarly, plant-derived hydrolysable biopolymers that presumably cycle more slowly^{26,28,36}, decreased by 14% between the 0 and 9 °C treatments, corresponding to $-0.65 \pm 0.2 \text{ mg g}^{-1}$ per °C increase in temperature ($r^2 = 0.52$, p < 0.0001; Fig. 1b, Fig. S2b), likely due to accelerated microbial decomposition of soil C^{4,37}. Surprisingly, the concentration of lignin phenols, which are made up of phenolic units that are expected to decompose faster under aerobic conditions^{8,9,15}, instead increased significantly in the surface peat by 12% between the 0 and 9 °C treatments (0.51 ± 0.2 mg g⁻¹ per °C warming; $r^2 = 0.18$, p = 0.01; Fig. 1d, Fig. S2d), indicating that warming might have induced lignin stabilization. Indeed, it is plausible that more lignin phenols entered soils via increased root litter production or that a shift in plant species composition toward shrubs richer in these compounds^{16,17} may explain this increase. It is also possible that this apparent increase in lignin phenols may be due to accelerated decomposition of more rapidly cycled labile C given that warming has increased the availability of labile sugars (and protein) in this experiment³⁸. In contrast, ratios of commonly used lignin oxidation (degradation) indices; acid-toaldehyde ratios of vanillyl and syringyl phenols, which typically serve as proxies for oxidation³², increased linearly with increasing temperatures (Fig. 2a-c), suggesting that warming, as expected, induced lignin degradation. The observed relative increase in lignin phenols (Fig.1d; Fig. S1) thus reflects a combination of increased organic matter inputs via plant litter and root exudates, enhanced degradation of more labile SOC molecules induced by higher temperatures^{16,38} and a better resilience of these molecules to warming as compared to the more labile ones.

Overall, the above findings indicate that warming stimulated a loss of SOC molecules, irrespective of their origins and potential decomposition rates - although lignin phenols showed a lower response than expected. These transformations likely reflect extensive decomposition driven by low water tables during summer dry periods¹⁷, compounded by warm temperatures and increased above- and belowground plant productivity^{16,17,20}, and labile C inputs to SOC^{38,39}. Warming is typically accompanied by a lower water-table and enhanced peat aeration, which can affect the temperature sensitivity of organic matter decomposition^{8,9,14,15}. Here, warming resulted in a substantial water level drawn down during summer dry periods (30 cm below hollow surface in warmer plots)¹⁷. Since soil microbial activity is intrinsically temperature sensitive^{5,40}, the lowered water table, together with additional plant litter and labile C inputs^{10,16,17,38} potentially stimulated microbial SOC transformation (priming), leading to measurable loss in SOC components.

We then assessed how warming combined with an elevated atmospheric CO₂ concentration (eCO₂) modified key SOC molecules. We tested the hypothesis that eCO₂ would result in increased plant litter production and C transfer from plant litter into SOC, thereby buffering warming-induced SOC losses as temperatures rise^{17,41,42}. Our results support this hypothesis and provide evidence that the interaction of warmer temperatures and eCO₂ led to increased plant C inputs to SOC in the surface peat, potentially offsetting the warming-induced C losses (Fig. 1; Fig. S2b). We observed that concentrations of hydrolysable biopolymers under eCO2 increased by 18% between the 0 and 9 °C treatments $(0.86 \pm 0.3 \text{ mg s}^{-1} \text{ per})$ °C warming; $r^2 = 0.42$, p < 0.0001; Fig. 1b; Fig. S2b) (indicating plant contribution), despite a non-significant increase in the concentration of solvent-extractable compounds (from $19.8 \pm 0.9 \text{ mg g}^{-1}$ in the +0 °C plot to $23.3 \pm 1.1 \text{ mg g}^{-1}$ in the +9 °C plot; *p* > 0.05; Fig. 1a). The source of hydrolysable biopolymers could be roots, as evidenced by elevated concentrations of root-specific suberin monomers (Fig. 3b) and root litter production¹⁶. Surprisingly, the concentration of lignin phenols decreased linearly with increasing temperatures

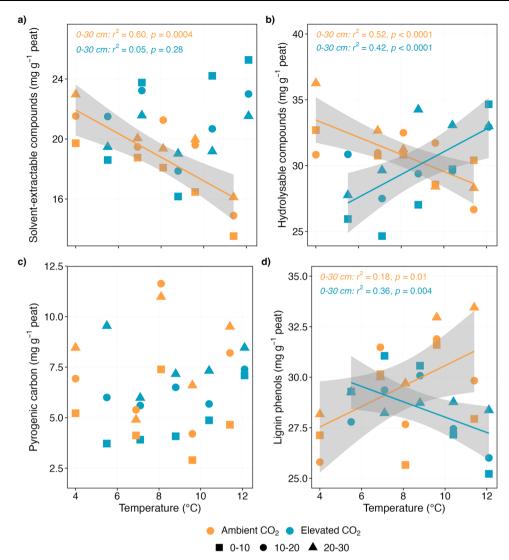


Fig. 1 | **The concentration of plant-, microorganism- and fire-derived SOC components.** Linear temperature response of the total sum of (**a**) solvent-extractable compounds (free lipids), (**b**) hydrolysable compounds (ester-bound lipids), (**c**) pyrogenic carbon and (**d**) lignin phenols in the surface peat (0–30 cm depth) following 4 years of warming and 2 years of elevated atmospheric CO₂ concentrations. The concentrations are plotted against average soil temperature measured at 0.3 m below the hollow surface from 2016 to 2018. Colours represent

ambient (orange) or elevated CO₂ (blue) treatment (n = 5 per treatment). Symbols represent different sampling depths. Lines indicate significant treatment effects p < 0.05. Linear regression with 95% confidence intervals is shown in grey. The absence of a line and/or confidence intervals indicates no significant trend. Note that the concentrations in the deeper anaerobic peat (>40 cm depth) did not differ significantly with increasing temperatures (see Supplementary Fig. 5).

 $(-0.38 \pm 0.2 \text{ mg g}^{-1} \text{ per }^{\circ}\text{C} \text{ warming}; r^2 = 0.36, p = 0.004; \text{ Fig. 1d}), \text{ rai-}$ sing possibilities of enhanced incorporation of other SOC compounds (i.e., hydrolysable biopolymers) into soil, leading to the decrease (dilution) of lignin phenols. Previous work at the study site demonstrated increased above- and belowground vascular plant biomass in response to warming and eCO₂ treatments^{16,17,20}. As the plant litter input increases, so do the concentrations of SOC components in our results (Fig. 1; Fig. S2b), which suggest that warming and eCO₂ could stimulate SOC incorporation via increased plant litter inputs^{17,42}, whereas general impacts on C storage remain unclear. However, it is plausible that our observed increase in SOC components under eCO₂ reflect a transient adjustment period to the 2 years of eCO₂ treatment, whereby plant and microbial growth and traits, and decomposition rates have not yet reached a new equilibrium⁴³. Alternatively, more rapid C cycling likely occurred within extant SOC components⁴¹, although an increase in SOC turnover appears to have been offset by additional plant productivity^{16,17}. Therefore, we posit that SOC will be partially preserved under warmer temperatures and greater CO_2 concentration in the short-term, though this may change in the future.

The above responses in SOC components to warming (only) and warming plus eCO_2 were much greater in the surface peat (0-30 cm) than at deeper depths (>30 cm; deep peat) (Figs. 1, 3; Fig. S5-6). We observed a rapid turnover of SOC molecules in the surface peat (Figs. 1-3), a high degree of transformation that likely reflects a combination of altered plant C inputs and decomposition driven by low water tables during summer dry periods, compounded by warmer temperatures (Fig. 4). Warming was accompanied by a substantial water level drawdown during summer dry periods (30 cm below hollow surface in warmer plots)¹⁷ in our study, and water table depth was correlated with SOC molecules (Fig. 4), suggesting that the warming and eCO₂ effect on SOC components was influenced by water level and downstream effects of peat oxygenation such as increased nutrient availability⁴⁴. However, it is worth noting that previous studies have not yet observed a direct effect of warming on soil moisture in this experiment¹⁷. Four years into the treatment, deep anaerobic peat

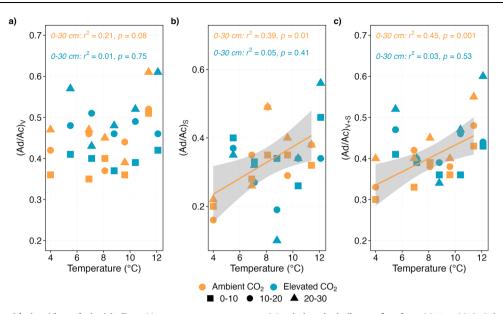
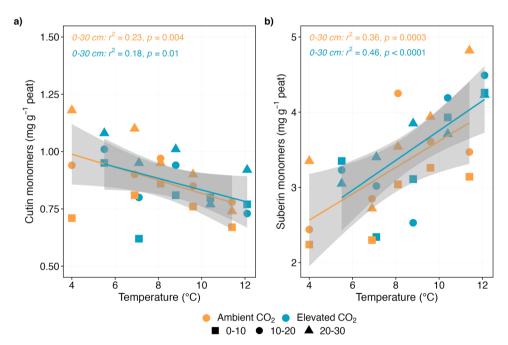
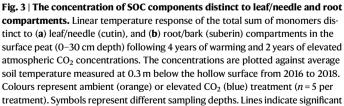


Fig. 2 | **Ratios of lignin oxidation (degradation) indices.** Linear temperature response of lignin oxidation proxy acid-to-aldehyde (Ad/Al) ratio for (**a**) vanillyl (V), (**b**) syringyl (S), and (**c**) vanillyl and syringyl phenols in the surface peat (0–30 cm depth) following 4 years of warming and 2 years of elevated atmospheric CO₂ concentrations. The ratios are plotted against average soil temperature measured

at 0.3 m below the hollow surface from 2016 to 2018. Colours represent ambient (orange) or elevated CO₂ (blue) treatment (n = 5 per treatment). Symbols represent different sampling depths. Lines indicate significant treatment effects p < 0.05. Linear regression with 95% confidence intervals is shown in grey. The absence of a line and/or confidence intervals indicates no significant trend.





treatment effects p < 0.05. Linear regression with 95% confidence intervals is shown in grey. The absence of a line and/or confidence intervals indicates no significant trend. Note that the concentrations in the deeper anaerobic peat (40–200 cm depth) did not differ significantly with increasing temperatures (see Supplementary Fig. 6). Biopolymers were selected according to their occurrence in the analysed leaves/needles, stems, and roots of the dominant plant species at SPRUCE (see Supplementary Table 2).

appears to be stable (Fig. S5-6) likely due to the slower decomposition that occurs under anoxic conditions^{2,9} and low-quality C substrates in these layers^{13,35}. It is also plausible that the deep anaerobic peat underwent extensive decomposition when it was part of the surface peat (1580-9200 BP)⁴⁵, creating peat that is less susceptible to the

changes that we observed in the surface layers. Together, our results suggest that warming and eCO_2 induce depth-dependent SOC responses, which should be carefully assessed to improve our mechanistic understanding and predictions of SOC dynamics under changing climate.

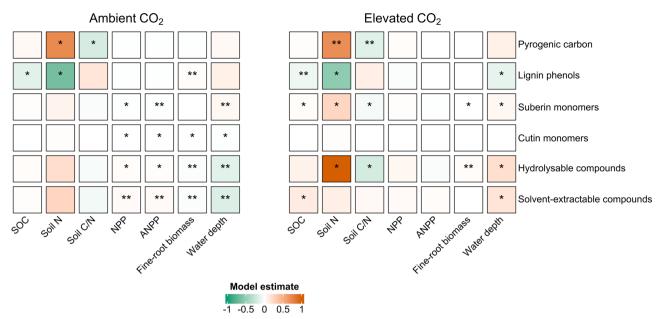


Fig. 4 | **Correlations** (*r*) between SOC molecules and biogeochemical drivers. Pearson correlations between the total sum of solvent-extractable compounds (free lipids), hydrolysable compounds (ester-bound lipids), cutin and suberin monomers, pyrogenic carbon, and lignin phenols (mg g peat⁻¹) and biogeochemical predictors soil organic carbon concentration (SOC; mg g peat⁻¹)^{17,43}, nitrogen concentration (soil N; mg g peat⁻¹)⁴³, carbon:nitrogen ratio (soil C/N; mg g peat⁻¹)⁴³, net primary productivity (NPP; g C m⁻² year⁻¹)¹⁷, aboveground net primary productivity (ANPP; g C m⁻² year⁻¹)¹⁷, fine-root biomass (g m⁻²)¹⁶, and maximum distance to the

Warming and eCO₂ enhanced the contribution of root-derived C inputs into SOC

To further delineate the contribution of plant litter inputs into SOC, we assessed changes in the incorporation and degradation of biopolymers distinct to either leaf/needle (cutin) or root/bark (suberin) compartments²⁶⁻²⁹ in the surface peat. The concentration of cutin monomers decreased linearly with increasing temperatures under both ambient CO2 and eCO2 (by ~20% between the 0 and 9 °C treatments, corresponding to -0.03 mg g^{-1} per °C warming) ($r^2 = 0.23$, p = 0.004 and $r^2 = 0.18$, p = 0.01 respectively; Fig. 3a; Fig. S3a). By contrast, the concentration of suberin monomers increased linearly with increasing temperatures in both ambient CO₂ and eCO₂ plots (by ~30% between the 0 and 9 °C treatments, corresponding to 0.19 mg g⁻¹ per °C warming) ($r^2 = 0.36$, p = 0.0003 and $r^2 = 0.46$, p < 0.0001respectively; Fig. 3b; Fig. S3b). Thus, warming and eCO₂ led to the loss of needle/leaf-derived C and gain of root-derived C in these soils (Fig. 3; Fig. S3). Importantly, root-derived C (inferred from suberin monomers) was positively correlated with SOC, fine root biomass and water table depth (Fig. 4) and fine root biomass significantly increased under warming and eCO2 (Table S1)¹⁶, implying increased root-derived C inputs within the surface peat. In corroboration of our results, litter manipulation experiments showed root-derived organic matter to be a source of SOC with greater relative stability and longer turn-over times²⁸, whereas leaf-derived C was found to be turned over more rapidly in (mineral) soils^{27,36,46}. Our study provides evidence that future climate change may lead to increased retention of root-derived inputs into SOC and a decline in leaf-derived C inputs in ecosystems such as boreal forested peatlands. As warming is likely to be accompanied by the decimation of peat-forming mosses and increases in fine-root biomass in peatland ecosystems^{16,47}, it is likely that root-driven accrual of SOC will represent an important mechanism for a continued C sequestration in these ecosystems in the short-term. However, the extent to which root-derived C can be incorporated and stabilized into deeper peat layers will depend on the depth at which root inputs water table (water depth; cm)^{17,47} in the surface peat (0–30 cm depth) following 4 years of warming and 2 years of elevated atmospheric CO₂ concentrations. Corrected significance at p < 0.05 is represented with *p < 0.01 is represented with ***, and p < 0.001 is represented with ***. The maximum distance (value) to the water table (relative to the hollows) was assessed from 2016 to 2018. The maximum values occurred in late summer (mid-August to mid-September). Additional descriptions of biogeochemical predictors and any data transformations are provided in the Methods.

increase. Thus, while we show that warming and eCO_2 , in the short term, enhanced incorporation of root-derived C into peat soils; it remains to be seen what the longer-term fate of this root-derived C will be.

There is no inherently stable peat SOC

We observed a rapid turnover of complex SOC molecules (such as lignin, cutin and suberin) and plant- and microbe-derived SOC components under warming-only and warming plus eCO₂ (Figs. 1, 3; Fig. S2), supporting the paradigm already observed in mineral soils that there is no inherently stable SOC^{24,25}. Complex SOC constituents such as lignin have traditionally been considered to cycle more slowly (on a multi-decadal time scale) relative to bulk SOC^{23,32}. However, the expectation that the enzymatic breakdown of complex SOC molecules is more sensitive to temperature in comparison to simple molecules³⁴, based in part on the kinetic theory, is not consistent with our results. We show rapid turnover of SOC molecules irrespective of their origins and potential decomposition rates (Fig. 1; Fig. S2). This high degree of transformation likely reflects extensive decomposition driven by low water tables¹⁷ and was compounded by warm temperatures along with increased labile organic matter³⁸ and plant inputs (Fig. 4). Our results are consistent with previous studies reporting a decline in aromatic compounds and soluble aromatics (indicative of lignin and pyrogenic C) in the water-extractable organic matter with warming and $eCO_2^{38,39}$. These results combined, show that microbial decomposition dynamics in this peatland are controlled by a range of biotic and abiotic ecosystem properties and not only by the kinetics of individual enzymatically catalysed reactions^{24,25}.

Unsurprisingly, molecules derived from historical pyrolysis (firederived carbon or pyrogenic C; PyC) were unaffected by warming or eCO_2 (Fig. 1c; Fig. S2c). We did not observe significant changes in PyC concentrations under ambient CO₂ or eCO_2 (p > 0.05; Fig. 1c; Fig. S4a), despite these peat soils containing PyC above global averages (-1.4% of the SOC; Fig. S1)⁴⁸. PyC did show an increasing trend with warming under eCO₂ (Fig. 1c; p > 0.05) probably due to previous biomass burning and incorporation of residues before the SPRUCE experiment⁴⁹. While these results are expected given that PyC is viewed as 'highly resistant organic matter^{23,30,32}, recent evidence suggests that the increased stability of these SOC components does not guarantee long-term persistence²³. To the best of our knowledge, no previous insitu peatland manipulation studies have reported on the fate of PyC under changing climate drivers.

Collectively, our study provides evidence that rising temperatures and atmospheric CO₂ levels will have cascading effects on the turnover rates and controls of SOC compounds that dictate peatland C storage. Specifically, climate-induced warming and associated water table drawdown may increase oxygen availability and accelerate microbial decomposition of SOC. We also highlight a rapid turnover of complex SOC molecules in response to warming and eCO₂, supporting the paradigm that SOC stability depends on ecosystem properties^{24,25}. Thus, SOC decomposition and turnover are predominantly determined by environmental and belowground plant-input constraints rather than temperature sensitivity of individual molecules²⁵. While our results indicate that ecosystem responses at SPRUCE are largely driven by surface peat, it remains to be seen whether a system shift characterized by more rapid C cycling to deeper depths in the peat column will be created by extended water-table drawdown and associated changes in plant and microbial community composition^{4,16,17,38} with further warming. Furthermore, we do not know whether the long-term SOC balance will be affected by plant-soil interactions or changes in soil microbial communities as they adapt to warmer temperatures⁵. Therefore, it is important to complement our results with long-term and time-resolved analyses. We also note that the results described here do not necessarily reflect the expected or observed responses for other peatland habitats. Nonetheless, the environmental gradients of SPRUCE provide us with model functions and parameters that can be applied to other similar peatland systems.

Implications

Understanding soil C dynamics, including vulnerability to warming and eCO₂, is critical for climate prediction over the coming decades to centuries^{5,7}. The unprecedented design of the SPRUCE wholeecosystem warming experiment enabled us to quantify changes in key SOC molecules to reveal mechanisms of climate change responses in boreal peatland C storage and cycling. We demonstrate that all SOC components responded to climate change. Under scenarios of water table drawdown, the resulting aerobic conditions and higher summer temperatures could cause substantial SOC loss, at least in the short term. Conversely, under a scenario of combined warmer temperatures and elevated atmospheric CO₂ concentration, enhance SOC storage is possible if C loss is balanced or exceeded by increased primary productivity, particularly from enhanced root growth, but this could be a short-term response. Moreover, warming could still cause considerable and rapid decomposition of previously water-saturated peat, which could make C stocks in high-latitude peatlands more susceptible to losses in the future. Peatlands build C stocks over centuries⁴⁵, but rising temperatures and atmospheric CO₂ concentrations rapidly changed the equilibrium at SPRUCE within a 4-year timescale, highlighting the vulnerability of these C-rich ecosystems to global change.

Methods

Site description

The Spruce and Peatland Response Under Changing Environments (SPRUCE) experiment is located at the southern boundary of the boreal region on the Marcell Experimental Forest in northern Minnesota, USA (47°30'20.5"N, 93°27'12.6"W; http://mnspruce.ornl.gov/). Mean annual temperature and precipitation is 3.4 °C and 780 mm respectively⁴⁹. The bog is ombrotrophic with peat depths of -3 m

(which accumulated over the last 11,000 years), with a pH ranging from 4.1 at the surface to 5.1 at 2 m depth^{45,49}. The bog is dominated by overstory trees, *Picea mariana* (black spruce) and *Larix laricina* (larch), and ericaceous shrubs, *Rhododendron groenlandicum* and *Chamaedaphne calyculata*, and bryophyte layer, primarily *Sphagnum* mosses. This ombrotrophic bog has a perched water table which fluctuates about 10–20 cm above the hollows after snowmelt, receding deeper later in the growing season^{17,45}.

Experimental design and sample collection

The SPRUCE experimental design is described in detail by Hanson, et al. ³³. Briefly, the experiment consists of ten octagonal transparent open-top enclosures of 12 m diameter and 7 m height. Whole-ecosystem warming is maintained at a series of increasing temperatures (regression design) to five levels (+0, +2.25, +4.5, +6.75 and +9 °C) down to a depth of -3 m using concentric arrays of electrical resistance heaters installed into peat and forced air warming. Peat warming was initiated between June and July 2014, following 3 months of a gradual treatment equilibration. Air warming was established in August 2015, thereby achieving whole ecosystem warming. In June 2016, eCO₂ treatment was introduced to the duplicate warming enclosures completing the planned experimental setup in this project. The eCO₂ treatment consists of elevating the local ambient atmospheric CO₂ concentration by +500 ppm (-900 ppm, with δ^{13} C-CO₂ isotope value of -54‰).

In August 2018, we randomly collected two soil cores from each of the plots in hollow microtopography where the surface of the hollow was defined as 0 cm. Surface samples (0-30 cm) were cut using a stainless-steel knife and extracted by hand, while a Russian peat corer was used to sample deeper peat (30-200 cm). Once collected, duplicate cores from the same plot were sectioned by depth (into 10 cm increments over 0 to 50 cm depth and 25 cm intervals from 50 to 200 cm), homogenized and combined to form a mixed sample. Dominant vascular plants and Sphagnum mosses were also randomly collected from each plot. Peat and plant samples were stored frozen at -20 °C immediately after sampling until further analyses. To avoid disturbance within the treatment plots, roots were collected from the dominant plants outside the enclosures and additionally picked from the soil manually. Peat sections and plant materials were later freeze-dried to constant weight. We sieved dried peat samples through a 5 mm mesh to remove larger litter fragments (2 mm sieve would have caused a loss of peat moss biasing the results). A subsample of the sieved peat and plant material was ground using a ball mill (MM400, Retsch, Haan, Germany) and then analysed for C concentration by an elemental analyser-isotope ratio mass spectrometer (EA-IRMS, Thermo Fisher Scientific, Bremen, Germany). Overall, SOC concentration increased with increasing depth from 44.0% at the surface to 52.5% at 2 m depth, but there was no effect of temperature or eCO₂ on the SOC concentration⁴³. Additionally, investigations of SOC stock at the site so far have shown that the mean effect of warming on soil C stocks was indistinguishable from zero¹⁷. Our observations are not contradictory with this result since the specific compounds we followed represent 20% of the total SOC (see Fig. S1) and should be seen as tracers of SOC. Bulk density measurements were taken from soil cores taken from each of the experimental plots in 2013 and 2020. Soil cores 5.2 cm in diameter were carefully excavated using a Russian corer and soil bulk density was calculated using the freezedried weights of the volumetric slices. Bulk density was used to estimate the mass of SOC molecular components (stocks) in the top 30 cm (g m⁻²). We calculated the mass of SOC molecular components by multiplying bulk density value (g soil cm⁻³) by concentrations of individual SOC compounds (solvent-extractable compounds, hydrolysable biopolymers, lignin phenols and pyrogenic C) (g g peat⁻¹) from each peat depth and summarized the values for the mentioned depth intervals.

Solvent-extractable compounds

Solvent-extractable lipids (alkanes, and alkanoic acids) in these soils have been previously reported⁴³. Here, we extend our previous study by including alkanols, steroids and terpenoids in our analysis in order to assess the stability of total solvent-extractable lipids (herein, solvent-extractable compounds) with varied degradability. These soils were analysed using the protocol reported in Wiesenberg and Gocke⁵⁰. Briefly, ~1g of plant materials and ~2g of milled peat was extracted using Soxhlet extraction with dichloromethane (DCM):methanol (MeOH) (93:7; v/v). The extracts were sequentially separated into low-polarity and acid (alkanoic acid) fractions using a KOH-coated (5%) silica gel column by eluting with DCM, and DCM:formic acid (99:1; v/v), consecutively. The low-polar fractions were further separated into aliphatic hydrocarbons (including alkanes, steranes and hopanes), polycyclic aromatic compounds, and heterocompounds (including alcohols, sterols and steranes) using an activated silica gel column by eluting with hexane, hexane:DCM (1:1; v/v) and DCM:MeOH (93:7; v/v), consecutively. An aliquot of the alkanoic acid fraction was spiked with an internal standard (eicosanoic acid; D₃₉C₂₀) and derivatized to fatty acid methyl esters (FAMEs) prior to analysis using boron trifluoride:MeOH solution (BF₃:MeOH). Aliphatic hydrocarbons, polycyclic aromatic compounds, and heterocompounds were spiked with tetracosane (D50C24), D10-phenanthrene and octadecanol (D37C18), respectively, as an internal standard prior to analysis. Overall, analytical errors were typically <10% based on replicate analysis (n = 6). Given that alkanes, alkanoic acids, alkanols, steroids and terpenoids showed similar responses to both warming and eCO₂, we summed them to represent solvent-extractable compounds.

Hydrolysable biopolymers

After pre-extraction of the solvent-extractable compounds, soil/plant residues were subjected to alkaline hydrolysis to extract ester-bound hydrolysable lipids (including cutin and suberin markers) as described by Mendez-Millan et al. ⁵¹. Solvent-extracted residues (-0.5 g of plant and -1 g of peat) were refluxed at 85 °C for 18 h with methanolic potassium hydroxide (KOH) (MeOH:water; 9:1, v/v; with 6% KOH). The extracts were filtered and acidified to pH 2.0 with 6 N hydrochloric acid (HCl). Compounds were recovered by liquid-liquid phase separation with DCM. The extracts were passed over sodium sulphate and dried under nitrogen gas. An aliquot of the extracts was spiked with an internal standard (eicosanoic acid; C₂₀) and derivatized with N,O-Bis(trimethylsilyl)-acetamide (BSA; Merck, Germany) for 1 h at 80 °C. Samples were analysed in duplicates and analytical errors were typically <10% (we report mean values n = 2).

Biopolymers distinct to either leaf/needle (cutin) or root/bark (suberin) were selected according to their occurrence in the analysed leaves, stems, and roots of the dominant plant species at SPRUCE; Picea mariana, Larix laricina, Rhododendron groenlandicum, Chamaedaphne calyculata, and Sphagnum mosses (Table S2). Leaves, stems, and roots were characterised by different abundances and chain lengths of *n*-alcohols. *n*-carboxylic acid. α -hydroxy alkanoic acid. ω -hydroxy alkanoic acid, α , ω -alkanedioic acid and mid-chainsubstituted hydroxy and epoxy alkanoic acids. The hexacosanoic acid (n-C26:0), octacosanoic acid (n-C28:0), dihydroxyoctadecanoic acid (9,10-diOH C_{18:0}) and dihydroxyoctadecenoic acid (x,18-diOH C_{18:1}) were used as markers for leaf/needle-derived C as they were either not detected or occurred in trace abundance in plant roots (Table S2). The ω -hydroxy alkanoic acids with a chain length of C₂₀, C₂₄ and C₂₆ and α,ω -alkanedioic acid with a chain length C₂₀, C₂₂ and C₂₄ (also α,ω octadecenoic acids, C18:1 diacid) were used as markers for root/barkderived C as they were either not detected or occurred in trace abundance in leaf/needle (Table S2). These compounds correspond to previously suggested cutin and/or suberin monomers^{27,51}. The unspecific monomers were treated as compounds derived from cutin or

Lignin phenols

Bulk peat samples (milled) were subjected to copper (II) oxide (CuO) oxidation to extract lignin phenols^{52,53}. Briefly, peat material equivalent to ~2 mg of organic carbon was mixed with 500 mg CuO. 50 mg of ammonium iron (II) sulphate hexahydrate and 20 mL of 2 M NaOH solution in Teflon-lined bombs. The bombs were then flushed with nitrogen for 10 min and heated (oxidized) in the microwave for 90 min at 150 °C. Once cooled, the oxidation products were spiked with ethylvanillin (3-ethoxy-4-hydroxy-benzaldehyde) and cinnamic acid (3phenyl-2-propenoic acid) as the recovery and internal standards, respectively, and subsequently acidified to pH 2.0 using 6 M HCl. The oxidation products were then extracted from the aqueous phase with ethyl acetate, dried under nitrogen and redissolved in p-Anisic acid. An aliquot of the oxidation products was converted to trimethylsilyl derivatives with N,O-bis (trimethylsilyl) trifluoroacetamide/ tetramethylchlorosilane (BSTFA + TCMS; 99:1; Sigma-Aldrich, USA) by heating for 20 min at 60 °C. Two analytical replicates were measured for all samples (analytical errors <20%; we report mean values n = 2). Eight characteristic oxidation products were summarized to represent the concentration of lignin phenols, including vanillyl (vanillin, acetovanillone, vanillic acid), syringyl (syringaldehyde, acetosyringone, syringic acid) and cinnamyl (p-coumaric acid, ferulic acid) phenols (Fig. S4b). In addition, CuO oxidation released salicylic acid, salicylaldehyde, and piceol phenols (see Fig. S4b for individual fractional abundances). Lignin-specific oxidation products were used to calculate lignin oxidation (degradation) indices; acid-to-aldehyde ratios of vanillyl and syringyl phenols³². We note that while the CuO method is not interfered by any other organic component in peat soils, the method may significantly reduce the yield of phenolic units upon CuO oxidation^{54,55}.

The above SOC molecular compounds were quantified using internal standards on gas chromatography (GC) equipped with a multimode injector and a flame ionization detector (GC-FID, Agilent 7890B). Compound identification was performed on an Agilent 6890 N GC equipped with split/splitless injector coupled to an Agilent 5973 mass selective detector (MS) using external standard mixtures and by interpretation of the mass spectra. On the GC-FID/MS, the samples (1µl) were injected in splitless mode. Both GC instruments were equipped with DB-5MS column (50 m x $0.2 \text{ mm} \times 0.33 \text{ }\mu\text{m}$), with helium as the carrier gas (1 ml min⁻¹). For solvent extractable compounds, details of the GC operating conditions are described elsewhere⁵⁰. For hydrolysable biopolymers, the GC oven temperature was held at 50 °C for 4 min, increased to 150 °C at a rate of 4 °C min⁻¹, then increased to 320 °C at a rate of 3 °C min⁻¹, with final isothermal hold at 320 °C for 40 min. Quantification of the biopolymers was achieved by comparison with an external calibration with 16-hydroxyhexadecanoic acid. For lignin phenols, the temperature was held at 80 °C for 5 min, increased from 80 °C to 110 °C at a rate of 2 °C min⁻¹, then increased to 170 °C at a rate of 0.5 °C min⁻¹, and at a rate of 15 °C min⁻¹ to 320 °C, with final isothermal hold at 320 °C for 10 min.

Pyrogenic carbon

Benzene polycarboxylic acids (BPCAs) were used as an approximation of pyrogenic carbon (PyC) and they were analysed following the protocol by Wiedemeier et al. ⁵⁶. Briefly, milled peat samples containing -10 mg of organic carbon were digested with 65% nitric acid in the digestion tubes at 170 °C for 8 h. The extracts were filtered and passed over a cation exchange resin for further cleaning and subsequently freeze-dried. The dried extracts were redissolved in MeOH:water (1:1; v/v) and passed over a solid phase extraction cartridge (C₁₈ SPE tube, Supelco, U.S.A), then dried under nitrogen and redissolved in

deionized water. Compounds were quantified on a high-performance liquid chromatograph (HPLC) (Agilent 1290 Infinity HPLC system, Santa Clara, U.S.A.), equipped with an Agilent InfinityLab Poroshell 120 SB-C₁₈ column (100 mm × 4.6 mm × 2.7 mm), and measured using a photodiode array detector (DAD). Ortho-Phosphoric acid (Honeywell, U.S.A.) dissolved in water (pH 1.2) was used as mobile phase A and pure acetonitrile (Scharlau, Spain) as mobile phase B. Samples were analysed in duplicates and analytical errors were typically <10% (we report mean values n = 2). All carboxyl functional groups (B3CA, B4CA, B5CA and B6CA) were summed to represent total concentration of PyC (see Fig. S4a for individual fractional abundances).

Data analyses

General linear mixed-effect models were used to determine the effects of temperature, elevated CO₂ concentrations and peat depth on individual SOC compounds (solvent-extractable compounds, hydrolysable biopolymers, lignin phenols and pyrogenic C). We checked for bivariate relationships between all variables to ensure that a linear model was appropriate. Normality and homoscedasticity in the models were checked using residuals and Q-Q plots (none of the data set was adjusted to fit parametric assumptions). In all cases, linear regression models included plot as a random effect, and all other predictor variables (temperature, elevated CO₂ concentrations and peat depth) as fixed effects. For instances in which both temperature and CO₂ effects were significant, we evaluated separate regressions against temperature for ambient and elevated CO₂ treatments (Table S3). When significant differences among depths were detected, we conducted pairwise comparisons using Tukey's honest significant difference test. In the models, we used the actual temperature measured at -0.3 m below the hollows averaged over the period 2016 to 2018 (Table S1). We chose 0.3 m as a representative depth because the water table has fluctuated about 30 cm relative to hollow during the growing seasons17.

Correlation heatmap (Pearson correlation) was calculated between individual SOC compounds (solvent-extractable compounds. hydrolysable biopolymers, lignin phenols and pyrogenic C) and possible predictors (SOC concentration, nitrogen concentration, carbon:nitrogen ratio, net primary productivity, aboveground net primary productivity, fine-root biomass, and maximum distance to the water table). Stepwise multiple linear regression with Akaike Information Criterion (AIC) as the model selection condition was used to investigate relationships between individual SOC compounds and the most important predictor variable(s). Only variables showing significant correlations with the investigated parameter (p < 0.1) were included as independent variables in the model. After the establishment of regressions, candidate models were carefully investigated for collinearity among the selected predictors and assumptions of normality by calculating variance inflation factors (a cut-off value of 1 when collinearity was considered not to exist) and graphically examining plots of residuals (log-transformation did not improve the overall distribution). In the multiple linear regression model, potential explanatory variables included: SOC concentration, nitrogen concentration, net primary productivity, fine-root biomass, and maximum distance to the water table (Fig. 4; Table S4).

Overall, the chosen level of significance was 5% (p < 0.05) in all statistical tests (unless stated otherwise). All data analyses were performed using the R v.4.2.0 (R Core Team, Vienna, Austria) using the RStudio interface v. 1.2.5033 (RStudio Team, PBC, Boston, MA).

Data availability

Data sets used in this study are available in the online SPRUCE project archive at https://doi.org/10.25581/spruce.113/2202278 and the longterm storage in the U.S. Department of Energy's Environmental Systems Science Data Infrastructure for a Virtual Ecosystem (ESS-DIVE; https://ess-dive.lbl.gov).

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Author contributions

M.W.I.S., N.O.E.O., A.M., G.L.B.W. and S.A. conceived the study; P.J.H. and C.M.I. designed and maintained the experiment; N.O.E.O., P.J.H., A.M., J.E.K., R.M.W. and C.M.I. carried out the field campaign; N.O.E.O. and G.L.B.W. carried out biomarker and geochemical laboratory analyses; R.M.W., P.J.H., N.O.E.O. and A.M. provided complementary data

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(including SOC and nitrogen concentration, net primary productivity, fine-root biomass, and water table depth); N.O.E.O. analysed the data with help from A.M. and S.A., and wrote the original draft. All authors commented on the interpretation and presentation of the data and contributed to editing the original draft. Funding acquisition: P.J.H., M.W.I.S. and N.O.E.O.

Competing interests

The authors declare no competing interests.

Additional information

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