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**LETTER**

Moderate nitrogen enrichment increases CO₂ sink strength in a coastal wetland

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E-mail: wangxy@igsnrr.ac.cn and [bxhie@yic.ac.cn](mailto:bhxie@yic.ac.cn)**Keywords:** nitrogen enrichment, coastal wetland, CO₂ sequestration, plant species composition, biomass distribution**Abstract**

Coastal wetlands remarkably influence terrestrial carbon (C) stock by serving as natural reservoirs for 'blue carbon'. Anthropogenic nitrogen (N) enrichment shapes the dynamics of soil and plant communities, consequently affecting the C balance and ecosystem functions. The impacts of various levels of N enrichment on CO₂ sequestration in coastal wetlands, however, remain elusive. Here we conducted a long-term field study of N fertilization in a coastal wetland in the Yellow River Delta, China, to investigate N effects on soil properties, indicators of plant dynamics, and fluxes of ecosystem CO₂. The results indicated that moderate N enrichment (5 g N m⁻² y⁻¹) stimulated C fluxes with increases in gross primary productivity (+26.4%), ecosystem respiration (+23.3%), and net ecosystem exchange (NEE, +31.5%) relative to the control. High (10 g N m⁻² y⁻¹) and extreme (20 g N m⁻² y⁻¹) amounts of N enrichment, however, had relatively minor impacts on these CO₂ fluxes. Overall, we observed a decrease in soil electrical conductivity (-24.6%) and increases in soil organic C (+25.2%) and microbial biomass C (+369.3%) for N enrichment. N enrichment also altered the composition of plant species, with a higher proportion of a local dominant species (*Phragmites australis*), and affected root biomass distribution, with more biomass near the soil surface. Structural equation modeling explained 65.2% of the variance of NEE and supported the assumption that N enrichment could alter the dynamics of soil properties and plant conditions and accelerate ecosystem CO₂ sequestration. These findings have important implications for forecasting the C cycle with increasing N deposition in coastal wetlands, contributing to the projections of the global C budget.

1. Introduction

Coastal wetlands are productive ecosystems where organic carbon (C) decomposes slowly in soil, so they function as reservoirs of 'blue carbon' and contribute as a natural climatic solution (Hopkinson *et al* 2012, Macreadie *et al* 2021). Recent studies on sinks of 'blue carbon' have emphasized the urgent need to restore and protect coastal wetlands due to their crucial role in global C sequestration (Han *et al* 2015). The influence of the increased deposition of nitrogen (N) on terrestrial C sinks, particularly in forests, has received

growing attention over the years (Reay *et al* 2008, Liu *et al* 2022). The effects and underlying mechanisms of N deposition on ecosystem C sequestration in coastal wetlands, however, remain subjects of ongoing debate. This uncertainty introduces noticeable uncertainties in global C budgets and model projections (Ward *et al* 2020).

Global annual N deposition rates have experienced a three- to fivefold increase over the past century, and projections suggest a further two- to threefold rise by the century's end (Lamarque *et al* 2005). The N and C cycles are intricately linked

(Gruber and Galloway 2008). Numerous studies have indicated that the effect of N deposition on the soil C sink relies on the interplay between two factors: (1) N-induced increases in soil C input of by stimulated plant growth and (2) N-induced losses of soil C from processes such as the decomposition of soil organic C (SOC), respiration, and mineralization (Reay *et al* 2008). For coastal wetlands that have been widely identified as N-limited ecosystems, long-term N enrichment could increase soil C respiration by altering the dynamics of soil, microbe, and plant communities (Macreadie *et al* 2021). Soil N availability could also regulate environmental responses to higher CO₂ levels, increased temperatures, and alterations in water availability, further influencing soil C dynamics (LeBauer and Treseder 2008, de Vries *et al* 2014, Du *et al* 2020). Plant growth and ecosystem function can be influenced by alterations in N cycling and soil quality caused by N deposition. Field studies support that N deposition could enhance plant growth and biomass accumulation in coastal wetlands (Guan *et al* 2019, Qu *et al* 2022), indicating the ability to take up more C with increasing N deposition. Increased inputs of soil C from aboveground litterfall and belowground root biomass could amplify soil C decomposition and release more C into the atmosphere. Increased N availability may also alter the soil environment (e.g. pH and electrical conductivity) and soil microbial activity and biomass, further affecting the levels and forms of N in coastal wetlands associated with soil heterotrophic respiration (Tao *et al* 2018). For example, moderate N addition could increase the rate of soil respiration, but extreme N input could lower soil pH, decrease microbial biomass, and decrease soil respiration (Wang *et al* 2019). Recent studies have also suggested that N addition could influence plant biomass distribution, regulating soil C cycling and balance (Li *et al* 2021, Keller *et al* 2023). The effects of N deposition on the sequestration and balance of ecosystem C are thus intricate and involve interactive plant and soil dynamics. Assessing the impact of N enrichment on C cycling and sequestration in coastal wetlands is challenging due to its combined direct and indirect effects on carbon cycle processes (Geoghegan *et al* 2018).

Our knowledge of the N effects on C cycling is growing, but we need an integrated understanding of its processes and mechanisms in coastal wetlands. We, therefore, conducted long-term field experiments in a coastal wetland of the Yellow River Delta (YRD), China. These experiments involved eight years (2012–2020) of N enrichment, subsequent with a campaign of measurements of soil properties, plant traits, and fluxes of ecosystem CO₂. The study had a dual purpose: firstly, to investigate the impacts of different levels of N enrichment on CO₂ sequestration in a coastal wetland, and secondly, to establish a connection between N-induced alterations in the soil

environment, plant dynamics, and the strength of the CO₂ sink in the wetland.

2. Methodology

2.1. N-fertilization experiment

The study was initiated in 2012 at the Ecological Research Station of Coastal Wetlands of YRD, Chinese Academy of Sciences (37°36'N, 118°57'E) (figure 1(a)). The YRD coastal wetlands are a relatively young and well-preserved ecosystem, with a rate of atmospheric N deposition of approximately 2.1 g N m⁻² y⁻¹, a mean annual temperature of 13 °C, and a total rainfall of 560 mm y⁻¹ (Guan *et al* 2019). The soil water salinity is relatively high due to the tide, ranging from 5 to 30 g l⁻¹ (Chu *et al* 2018). The dominant herbaceous species in the area are *Phragmites australis* and *Suaeda salsa*, with average heights of ~1.7 m (Han *et al* 2015).

We established 20 experimental plots (1 m × 1 m) in an expansive field (50 plots in total), which were sparsely distributed to prevent the potential movement and loss of N (figure 1(b)). We tested four rates of N fertilization (N0: 0 g N m⁻² y⁻¹, N5: 5 g N m⁻² y⁻¹, N10: 10 g N m⁻² y⁻¹, and N20: 20 g N m⁻² y⁻¹), with each rate having five replicates, for a total of 20 plots. The selected fertilization rates were aimed at replicating levels of anthropogenic atmospheric nitrogen deposition ranging from normal to severe. Specifically, these rates were used to simulate normal (N0), moderate (N5), high (N10), and extremely high (N20) rates of atmospheric N deposition (Guan *et al* 2019, Wilcots *et al* 2022). We evenly sprayed ammonium nitrate (NH₄NO₃), a common fertilizer (Ahlgren *et al* 2008) dissolved in 1.2 l of deionized water, once a month to the fertilized plots since 2012. For N0 control plots, we applied deionized water without fertilizer to ensure a consistent soil-water content. To prevent the plants from intercepting the fertilizer, we applied the fertilizer directly onto the soil or the roots of the plants through spraying.

2.2. Measurement of fluxes of ecosystem CO₂

To measure fluxes of ecosystem CO₂, we used an Ultraportable Greenhouse Gas Analyzer (U-GGA 915, GLA132 Series, Los Gatos Research Inc., San Jose, USA) attached to a transparent chamber (diameter: 0.4 m; height: 1 m) (figure 1(c)). We inserted one circular frame (diameter: 0.4 m; height: 0.1 m) into the soil in each plot at a depth of 7 cm in July 2018. We kept two small electric fans operating continuously during the measurements to ensure proper air mixing inside the chamber. Additionally, when the plants exceeded a height of 1 m, we introduced an additional chamber measuring (diameter: 0.4 m; height: 0.7 m) for the measurements. After the gas concentration in the chamber began to increase linearly, nine

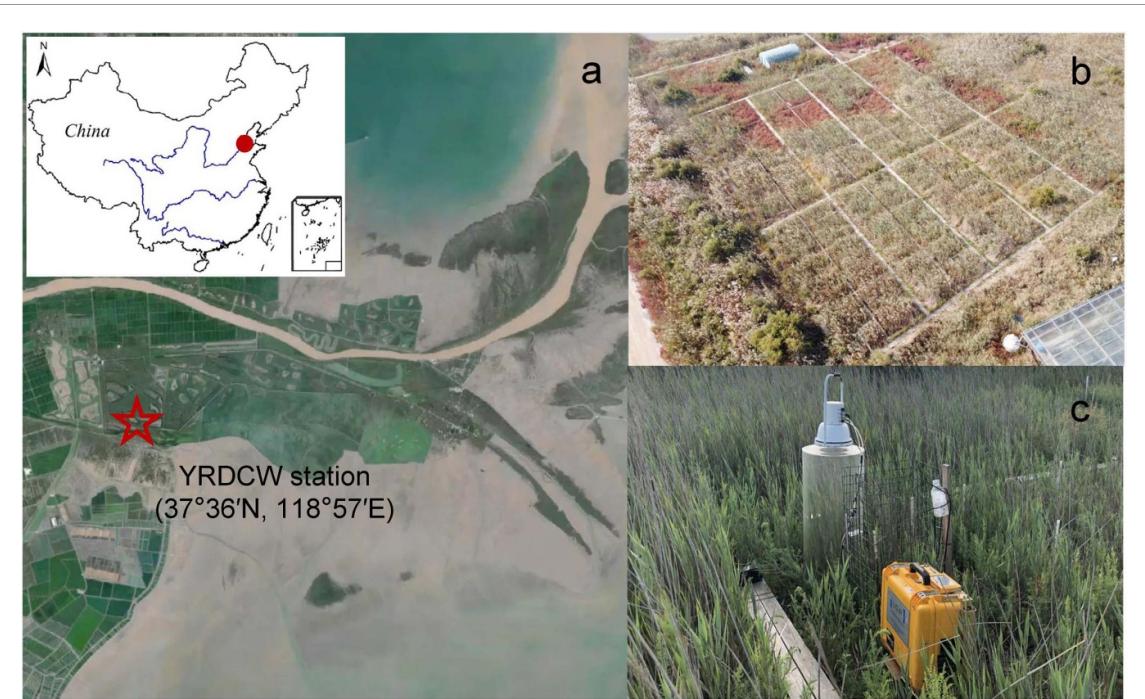


Figure 1. Descriptions of field experiment. (a) The location of the Yellow River Delta Coastal Wetlands (YRDCW) station with the corresponding latitude and longitude. The background is a high-resolution satellite image of YRD. (b) Bird view of the experimental field taken by a drone. (c) The equipment used to measure the fluxes of ecosystem CO₂.

CO₂ concentrations were measured in succession at 10 s intervals (Sun *et al* 2022). NEE was determined by calculating the rate of change in CO₂ concentration over time. Subsequently, the chamber was ventilated, repositioned on the frame, and covered with an opaque cloth to measure ER. In our study, GPP was determined as the difference of ER and NEE. It should be noted that positive values of GPP and ER represent opposite directions of C fluxes and negative (positive) NEE suggests an ecosystem C sink (source).

To examine the effects of eight-year N enrichment on C fluxes, we conducted two-year continuous measurements of CO₂ fluxes. Ecosystem CO₂ fluxes were measured twice a month on clear and sunny days at noon (around 12:00) from 1 January 2019 to 31 December 2020. The mid-day GPP and NEE represent the daily peak of corresponding CO₂ fluxes, to some degree, amplifying the N signal of CO₂ sink strength (Xia *et al* 2009, Sun *et al* 2022). It should be noted that the scaling of CO₂ fluxes that converts once-a-day values to daily total or averaged values is only possible with hourly continuous measurements due to varying diurnal patterns of CO₂ fluxes throughout different seasons (Reichstein *et al* 2005). In our study, we only focused on the directly measured dynamics of mid-day CO₂ fluxes under N enrichment, excluding the uncertainties caused by the scaling processes.

2.3. Measurement of plant indicators

We measured multiple plant traits including aboveground and root biomass, total number of plants, and the leaf area index (LAI). We applied an ACCUPAR LP-80 ceptometer (METER Group, Pullman, USA) to measure LAI in August 2020. We evenly divided each plot (1 m × 1 m) into four quadrants using a cross at the end of the growing season. Then we randomly selected a quadrant, which represents 1/4 of the total area (0.25 m × 0.25 m), to determine total number of plants and percentage of each plant species, and to collect plant aboveground biomass. For the sampling, we trimmed the aboveground vegetation to ground level, removed standing dead material, and classified the vegetation by type. The species composition was determined as the percentage of plant number for each species we sampled. For root sampling, we used a root auger (diameter of 10 cm) to take root samples, and then calculated the root biomass of each sample plot based on the area ratio. We collected the roots at multiple depths to measure their biomasses: 0–10, 10–20, 20–30, and 30–40 cm. The roots were divided into coarse and fine roots, defined by diameters ≥ 2 and < 2 mm, respectively. We placed the plant aboveground and root samples into sturdy envelopes and subjected them to a 48 hour drying process in an oven set at 80 °C to measure aboveground biomass and root biomass (g m⁻²).

2.4. Measurement of soil properties and microbes

We collected soil samples (20 cm depth) and measured data for soil properties in October 2020 in five randomly selected experimental plots (1 m × 1 m) in July 2020. The soil samples were air-dried for measuring soil properties. We measured extractable N ($\mu\text{g g}^{-1}$ dry soil) in the soil samples, i.e. the total amount of NH_4^+ -N and NO_3^- -N, by mixing 3 g of soil (dry mass) with 15 ml of 2 M potassium chloride (KCl). Soil pH was potentiometrically determined in a 1:5 (w/v) soil: water suspension using a portable pH meter. We employed three parameter sensors for soil electrical conductivity measurement (EC, ds cm^{-1}) in the top 10 cm of soil. The soil samples for SOC analysis were pretreated with hydrochloric acid (1 mol l^{-1}) to remove inorganic C. The samples were subsequently analyzed using the elemental analyzer (Vario MACRO cube, Elementar Analysensysteme, Germany).

We used soil samples (fresh soil) to determine microbial biomass C (MBC) and microbial community structure under different N treatments. In our study, the measurement of MBC was carried out using a Shimadzu TOC analyzer (TOC-VCPh) along with the chloroform-fumigation extraction method (Brookes *et al* 1985). To examine the microbial community structure, we utilized the PowerSoil DNA Isolation Kit (MO BIO Laboratories, Inc., USA) for the extraction of total DNA from the samples, which were subsequently stored at -20°C . In order to identify functional microorganisms, the bacterial 16 S rRNA gene in V3-V4 fragments was targeted using the primers 341F and 806R (Mori *et al* 2014). The reaction setup consisted of a total volume of 50 μl , including 25 μl of $2 \times$ Primer Taq (TaKaRa, RR902A), 1 μl of forward primer (10 mM), 1 μl of reverse primer (10 mM), 3 μl of g-DNA, and 20 μl of nuclease-free water. The thermal cycling protocol involved the following steps: (1) an initial denaturation at 94°C for 5 min, (2) 31 cycles of denaturation at 94°C for 30 s, (3) annealing at 52°C for 30 s, (4) extension at 72°C for 45 s, (5) a final extension at 72°C for 10 min, and (6) a hold at 72°C before purification (Xu *et al* 2016). Gel electrophoresis was employed to analyze the PCR products, while pyrosequencing was performed on the MiSeq platform. To ensure data quality, Trimmomatic-0.33 was utilized for data filtering. Additionally, the MOTHUR project (Li *et al* 2022) was used to identify and remove any reads suspected to be chimaeras. Operational taxonomic units (OTUs) at the species level were generated by setting a threshold of $\geq 97\%$ to quantify the similarity between sequences (Xu *et al* 2016, Gu *et al* 2019). Taxonomic assignments were completed using the UCLUST method with the UNOISE3

algorithm for error-correction, adopting a similarity threshold of 0.5. Finally, reference sequences were obtained from the Greengenes OTU database (gg_13_8_ots, http://qiime.org/home_static/dataFiles.html). We measured the soil bacterial compositions for topsoil (0–20 cm) and subsoil (20–40 cm).

2.5. Statistical analyses

The N level was the only controlling factor in our experiment, so we performed a one-way analysis of variance (ANOVA) using IBM SPSS Statistics (www.ibm.com/spss). Statistical significance was determined at $p < 0.05$. We used Pearson correlation analyses to assess the associations among the soil properties, plant growth, species composition, biomass distribution, and CO_2 fluxes. We used structural equation modeling (SEM) to examine the underlying processes and mechanisms to quantitatively elucidate the direct and indirect effects of N enrichment on ecosystem C sequestration (i.e. NEE). We assumed that N enrichment regulated NEE by affecting the soil properties, plant growth, species composition, and distribution of root biomass. We excluded the extreme N treatment (N20) from the SEM to better examine the responses of NEE to N enrichment. To assess the goodness-of-fit of the SEM, we calculated several metrics outlined by Hair *et al* (2017): the Goodness-of-Fit Index (GFI), the standardized root mean square residual (SRMR), the ratio of χ^2 to the degrees of freedom, and the root mean square residual (RMR). Due to the limited sample size (< 100), we utilized partial least squares in the SEM analyses using the 'Lavaan' package in R (v.4.1.1) (Rosseel 2012). In our study, we measured two-year C fluxes (from January 2019 to December 2020) and one-year plant-soil indicators (2020). Our main analyses were based on measurements in 2020 to avoid potential uncertainty caused by unmatched time frame.

3. Results

3.1. Impacts of long-term N enrichment on the fluxes of ecosystem CO_2

The temporal patterns of the CO_2 fluxes (i.e. NEE, ER, and GPP) in the N treatments were similar in 2019 and 2020 (figures 2(a), (c), and (e)), with an upward (downward) trajectory followed by a peak and subsequent decrease (increase). We detected considerable discrepancies, especially in the rainy season (from 15th July to 15th September). Moderate N enrichment (N5) stimulated CO_2 fluxes with increases in NEE (+31.5%, $p < 0.05$), ER (+23.3%, non-significant) and GPP (+26.4%, $p < 0.05$) throughout the full year, growing season, and rainy season

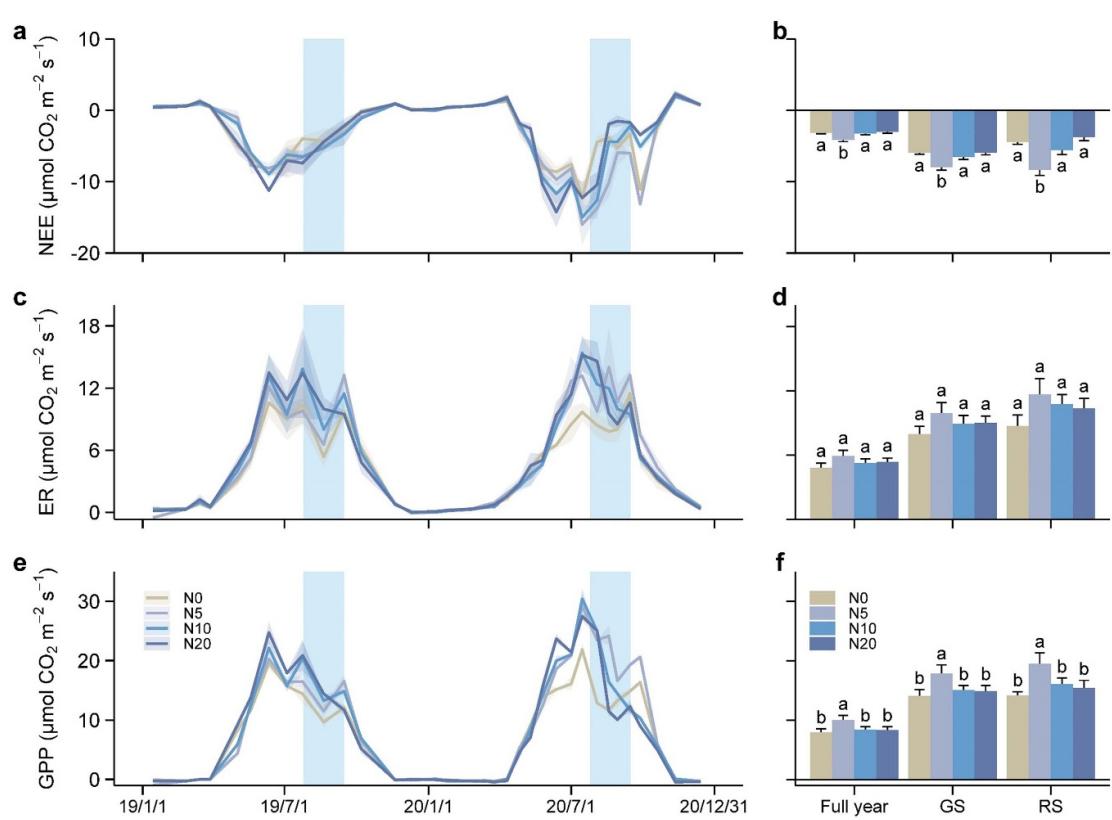


Figure 2. Effects of long-term N enrichment on the fluxes of ecosystem CO_2 . (a), (c), and (e). Variations of net ecosystem exchange (NEE), ecosystem respiration (ER), and gross primary productivity (GPP) (means \pm standard deviations) for 2019 and 2020, respectively. The light-blue bars represent the rainy season (RS, from 15th July to 15th September). (b), (d), and (f). Measurements for the entire year, the growing season (GS, NEE < 0), and the RS. The N-enrichment treatments, denoted as N0, N5, N10, and N20, correspond to the application of 0, 5, 10, and $20 \text{ g N m}^{-2} \text{ y}^{-1}$, respectively. Different letters represent significant differences ($p < 0.05$).

(figures 2(b), (d), and (f)). High (N10) and extremely high (N20) N enrichment had relatively minor effects on ecosystem CO_2 fluxes.

3.2. Responses of plant dynamics to long-term N enrichment

Aboveground biomass was generally higher with N enrichment, 45.4% higher in N10 than N0 (figure 3(a)). The number of plants was slightly higher in N5 (+21.4%) and N10 (+21.3%) and slightly lower (-8.1%) in N20 relative to N0 (figure 3(c)). The impact of N enrichment on LAI was not obvious, but LAI was highest in N5 (figure 3(d)). The *P. australis* and *S. salsa* were dominant species over the fields (figure 3(b)). The ratio of these two dominant species showed a significant difference ($p < 0.05$) in species composition under N5 (+14.5%) and N10 (+12.8%) relative to N0 (figure 3(e)). The biomasses of coarse and fine roots were significantly higher ($p < 0.05$) by 126.7% and 50.8%, respectively, in N10 than in N0 (figure 4(a)). The ratio of coarse-root to fine-root biomass was slightly higher under N enrichment (figure 4(b)). N enrichment relocated more of the coarse-root biomass into near-surface soil (figure 4(c)) but had a

minor impact on the distribution of fine-root biomass (figure 4(d)).

3.3. Effects of long-term N enrichment on soil properties and microbes

The concentration of extractable N was significantly higher ($p < 0.05$) in N5, N10, and N20 than in N0 by 71.8, 197.7, and 371.4%, respectively (figure 5(a)). Soil pH did not differ significantly among the N treatments (figure 5(b)). EC was significantly lower ($p < 0.05$) in N5, N10, and N20 than in N0 by 13.8, 29.4, and 30.5%, respectively (figure 5(c)). SOC concentration was significantly higher ($p < 0.05$) in N5 (22.5%), N10 (33.9%), and N20 (19.9%) compared to N0 (figure 5(d)). We found apparent increases in MBC under N enrichment, with 378.5%, 301.9%, and 427.5% increases in N5, N10, and N20, respectively, compared to N0 ($p < 0.05$) (figure 6(a)). Soil bacterial compositions and corresponding percentages were overall similar under different N treatments in topsoil (0–20 cm) and subsoil (20–40 cm) (figures 6(b) and (c)), with relatively higher proportions for *Proteobacteria* (topsoil: $29.7 \pm 1.7\%$, subsoil: $34.2 \pm 6.8\%$) and *Chloroflexi* (topsoil: $20.7 \pm 0.7\%$, subsoil: $15.5 \pm 3.3\%$).

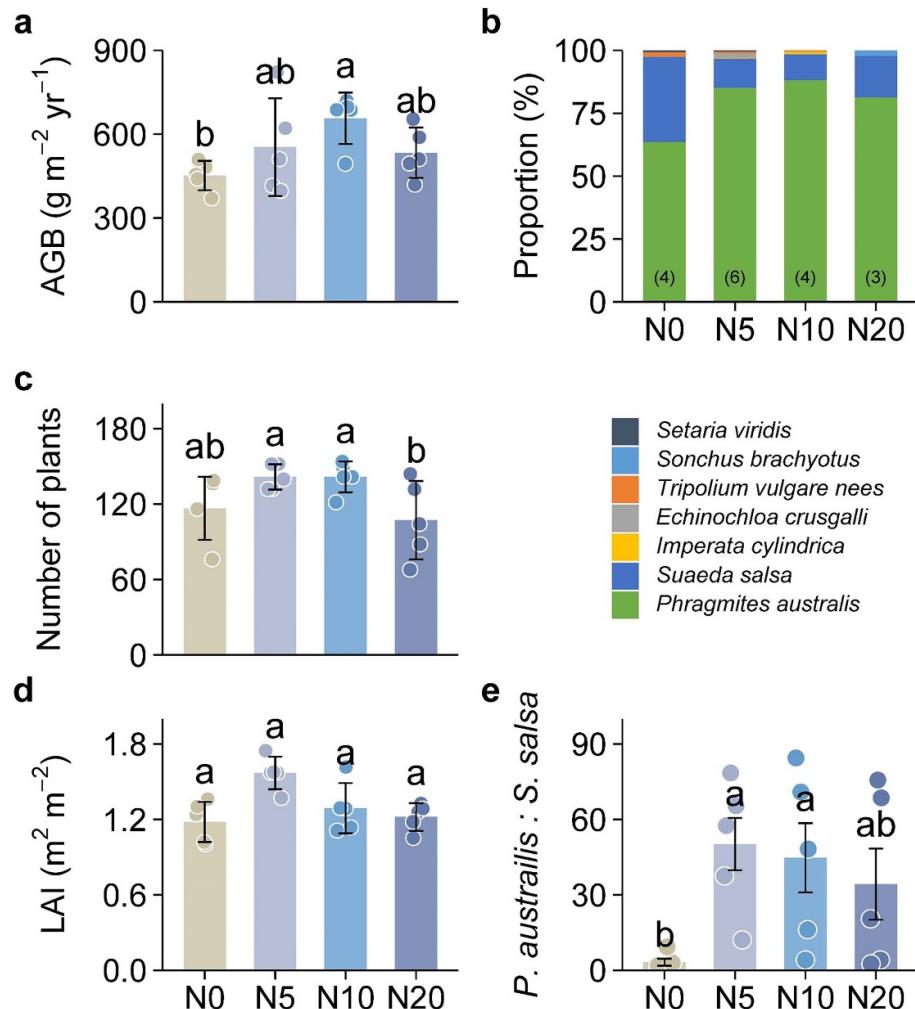


Figure 3. Effects of long-term N enrichment on plant growth and species composition. (a), (c), and (d). Distributions of aboveground biomass (AGB), number of plants, and the leaf area index (LAI) (means \pm standard deviations), respectively. (b). Species composition. The numbers in the columns represent the total number of species (i.e. species richness) in the treatments. (e) Ratio of the two dominant plant species (*P. australis* and *S. salsa*). The N-enrichment treatments, denoted as N0, N5, N10, and N20, correspond to the application of 0, 5, 10, and 20 g N m⁻² y⁻¹, respectively. The dots represent the value of replicated observations ($n = 5$). Different letters represent significant differences ($p < 0.05$).

3.4. Mechanisms of N effects on ecosystem C sink in the coastal wetland

Extractable-N concentration was correlated negatively with EC ($r = -0.51$; $p < 0.01$) and positively with SOC concentration ($r = 0.84$, $p < 0.001$), MBC concentration ($r = 0.63$, $p < 0.01$), above-ground biomass ($r = 0.62$, $p < 0.01$), number of plants ($r = 0.55$; $p < 0.05$), root biomass ($r = 0.77$; $p < 0.01$), and the biomass ratio between topsoil (0–20 cm) and subsoil (20–40 cm) ($r = 0.56$; $p < 0.01$) (figure 7). Extractable-N concentration was not significantly correlated with the CO₂ fluxes. The indicators of plant growth (aboveground biomass, number of plants, and LAI), however, were negatively correlated with NEE ($p < 0.05$). The SEM analysis generally confirmed the negative effects of N enrichment on NEE (figure 8). The criteria of the significance of the fitting of the SEM model were satisfied, with $\chi^2 = 8.433$, $p = 0.49$ (>0.05), GFI = 0.94 (≥ 0.9), SRMR = 0.07 (≤ 0.08), and RMR = 0.06 (≤ 0.08).

The fitted model explained 65.2% of the variance of NEE. Specifically, higher N availability could directly increase plant growth (i.e. plant number and above-ground biomass) and then decrease NEE. N enrichment could also reduce EC and further increase the ratio of coarse to fine roots and root biomass, leading to negative and positive effects on NEE, respectively.

4. Discussion

4.1. Changes in soil properties and plant dynamics induced by N enrichment

The coastal wetlands in YRD experience periodic tides that cause them to alternately submerge and become exposed, leading to the accumulation of salt and leaching with high EC (Han *et al* 2015). This environmental stressor could influence soil microbial activity and the cycling of soil C (Chambers *et al* 2011). We found that eight-year N enrichment substantially reduced EC (figure 5(c)), consistent with

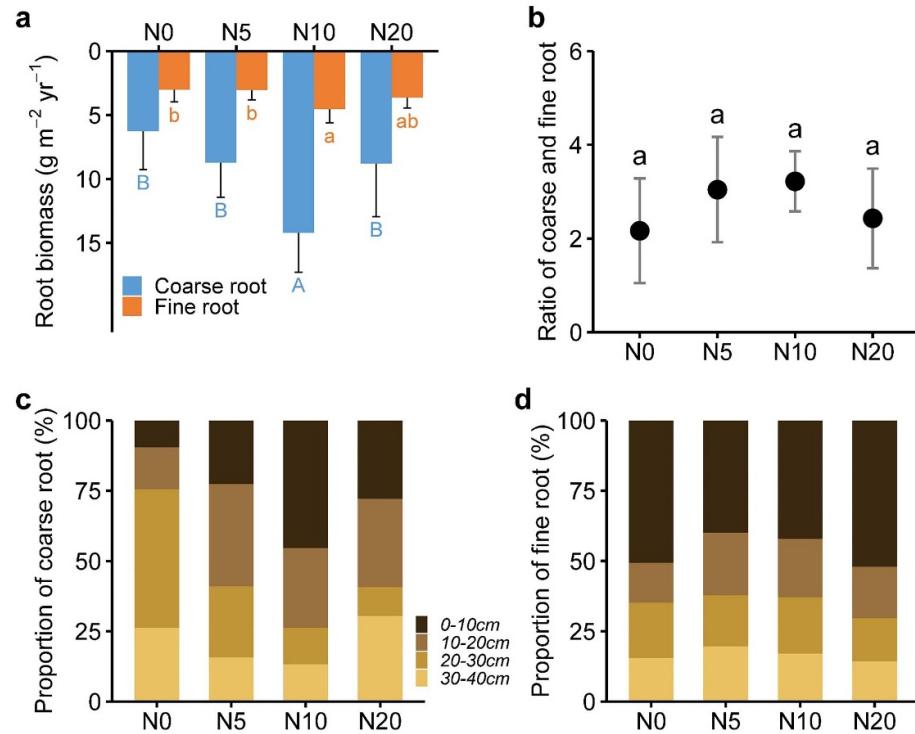


Figure 4. Effects of long-term N enrichment on the distribution of root biomass. (a) and (b) Distribution of root biomass and the ratio of coarse to fine roots (means \pm standard deviations), respectively. (c) and (d) Distribution of biomass in gradients of soil depth (0–40 cm) for coarse and fine roots, respectively. The N-enrichment treatments, denoted as N0, N5, N10, and N20, correspond to the application of 0, 5, 10, and 20 g N m⁻² y⁻¹, respectively. Different lower- and uppercase letters represent significant differences ($p < 0.05$).

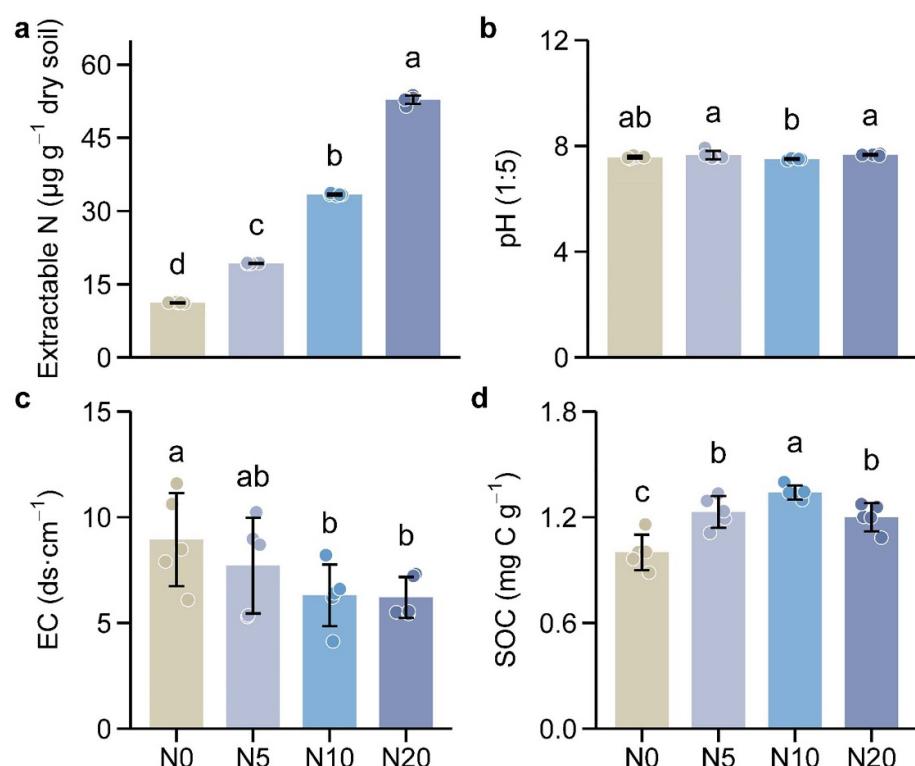


Figure 5. Effects of long-term N enrichment on soil physiochemical properties. (a)–(d). Distributions of soil extractable N ($\mu\text{g g}^{-1}$ dry soil), pH, EC (ds cm^{-1}), and soil organic C (SOC) with N addition (means \pm standard deviations), respectively. The N-enrichment treatments, denoted as N0, N5, N10, and N20, correspond to the application of 0, 5, 10, and 20 g N m⁻² y⁻¹, respectively. The dots represent the value of replicated observations ($n = 5$). Different letters represent significant differences ($p < 0.05$).

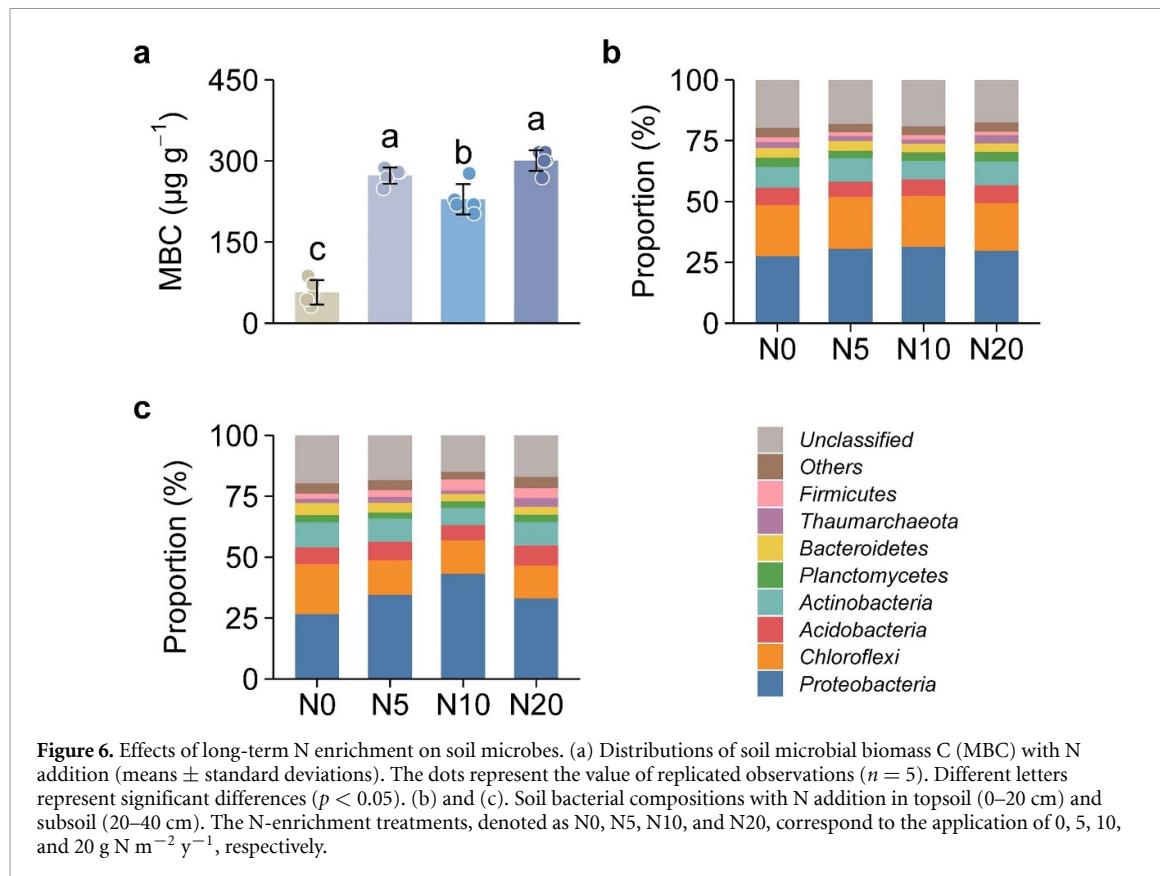


Figure 6. Effects of long-term N enrichment on soil microbes. (a) Distributions of soil microbial biomass C (MBC) with N addition (means \pm standard deviations). The dots represent the value of replicated observations ($n = 5$). Different letters represent significant differences ($p < 0.05$). (b) and (c). Soil bacterial compositions with N addition in topsoil (0–20 cm) and subsoil (20–40 cm). The N-enrichment treatments, denoted as N0, N5, N10, and N20, correspond to the application of 0, 5, 10, and 20 g N m $^{-2}$ y $^{-1}$, respectively.

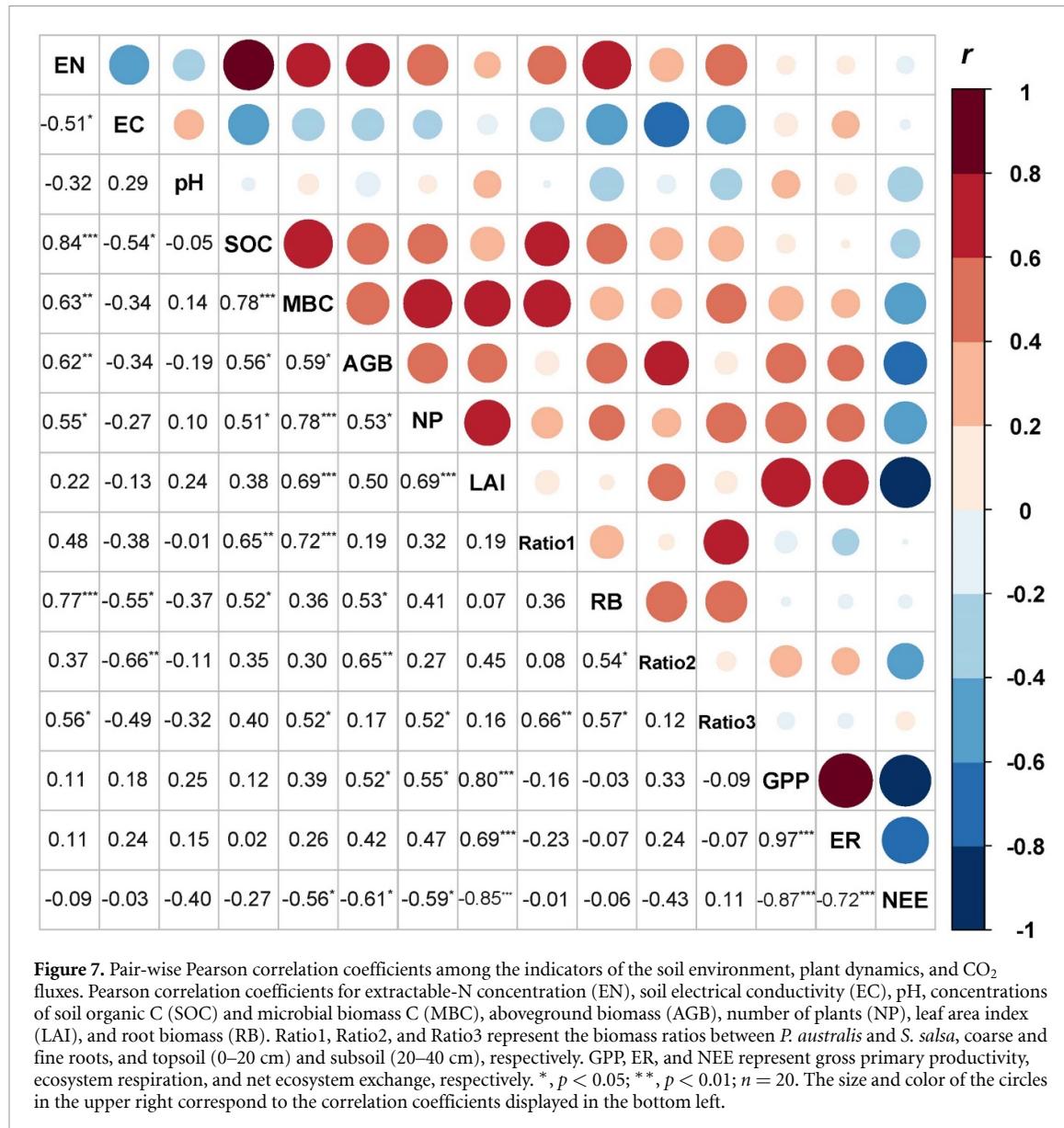
previous studies (Guan *et al* 2019, Qu *et al* 2019). Increases in the availability of soil N potentially alleviate salinity stress in coastal wetlands, which is beneficial to soil microbes and plant communities (Qu *et al* 2020, Huang *et al* 2022). SOC concentration can be a proxy of the strength of sinks of soil C, and the SOC concentration in our study was significantly higher under N enrichment (figure 5(d)), supporting a stronger C sink controlled by the availability of N. The N-induced increase in MBC helps explain the higher ER under N enrichment (figure 6(a)), to some extent, offsetting the C sink strength in coastal wetlands. Due to the tidal erosion, only salt-tolerant microbes can survive in the coastal wetlands of YRD (Kearns *et al* 2016, Li *et al* 2022), leading to a stable diversity of microbes even with N addition (figures 6(b) and (c)).

So far, we have relatively limited knowledge of the effects of N on plant biophysical dynamics in coastal wetlands. Long-term N enrichment strongly regulated the species composition of the plants in the YRD coastal wetland, with an increase in *P. australis* and a decrease in *S. salsa* (figure 3(e)). This alteration indicated a species-dependent adaption and suitability under anthropogenic nutrient enrichment (Porter *et al* 2013). Changes in species composition caused by N deposition have the potential to modify microclimates, sources of organic matter, and biogeochemical processes (Hu *et al* 2020). Root traits represent the primary source of input of SOC and C stocks

in local ecosystems (Mokany *et al* 2006). Increased N availability in our study stimulated root biomass (figure 5(a)), confirming the nutrient stress of the plant community in this YRD coastal wetland (Guan *et al* 2019). Interestingly, the biomass of coarse roots was relocated from the subsoil to the near-surface soil (figure 4(c)), suggesting the influence of N stress on root production in coastal wetlands (McMurtrie *et al* 2012).

4.2. Effects of long-term N enrichment on the sequestration of ecosystem CO₂ in the coastal wetland

We found that moderate N enrichment increased the CO₂ fluxes in coastal wetlands. Excessive N input, however, can partially or fully offset these effects, consistent with N saturation previously reported (Wei *et al* 2012, de Vries *et al* 2014). Based on our correlations and SEM analyses, we found a potential link between N enrichment and C sink associated with dynamics of soil and plants (figure 8). Our study generally supported the N-induced increase in CO₂ release (Sun *et al* 2023), with a relatively high ER with large increase in MBC under N enrichment (figures 2(c) and 6(a)). More importantly, the increased uptake of CO₂ by plants by photosynthesis (GPP) was higher than the increased release of CO₂ by ER under N enrichment that led to a higher NEE, suggesting the potential of a stronger CO₂ sink under future scenarios with increasing N



deposition (Liu *et al* 2005). The positive impacts of N deposition on plant growth and ecosystem C sink in coastal wetlands could also be controlled by N availability, supported by previous studies (LeBauer and Treseder 2008, Johnson *et al* 2016, Li *et al* 2022). We found a lower response of NEE and GPP to extremely high N deposition than to moderate N deposition (figure 2), suggesting a potential tipping point of N deposition effect on ecosystem C sequestration in coastal wetlands. Divergent magnitudes and patterns of C fluxes across wetlands were widely reported, suggesting a high spatial heterogeneity of environmental and climatic responses of C fluxes (Lu *et al* 2017, Hu *et al* 2020). For example, CO_2 emissions (ecosystem respiration) in 21 coastal wetlands ($87.64 \text{ mg} \cdot \text{m}^{-2} \cdot \text{h}^{-1}$) were much higher than that in inland wetlands ($39.17 \text{ mg} \cdot \text{m}^{-2} \cdot \text{h}^{-1}$). Even within coastal wetlands, large differences in C fluxes in China and North America were reported (Hu *et al* 2020), highlighting the discrepancy of driving factors.

4.3. Limitations and implications

Limited field experiments inhibit our understanding of the effects of N deposition on CO_2 cycling and ecosystem functions in coastal wetlands, unlike intensively investigated forests. We conducted a field study using eight-year N fertilization to investigate how N enrichment affected CO_2 sequestration, considering relevant biotic and abiotic factors. Our study however, still contained limitations of study design, assumptions and constraints. First, we only measured CO_2 fluxes for two years to analyze the effects of N for only one site. This short study period and limited spatial scale may cause some uncertainty if other environmental factors, e.g. air temperature and precipitation, may jointly influence the effects (Greaver *et al* 2016). Second, limited observations especially in nighttime may influence the responses of C fluxes to N deposition. More frequent measurements (hourly) are needed to reduce the uncertainty and enhance the understanding of N enrichment on daily cycles of

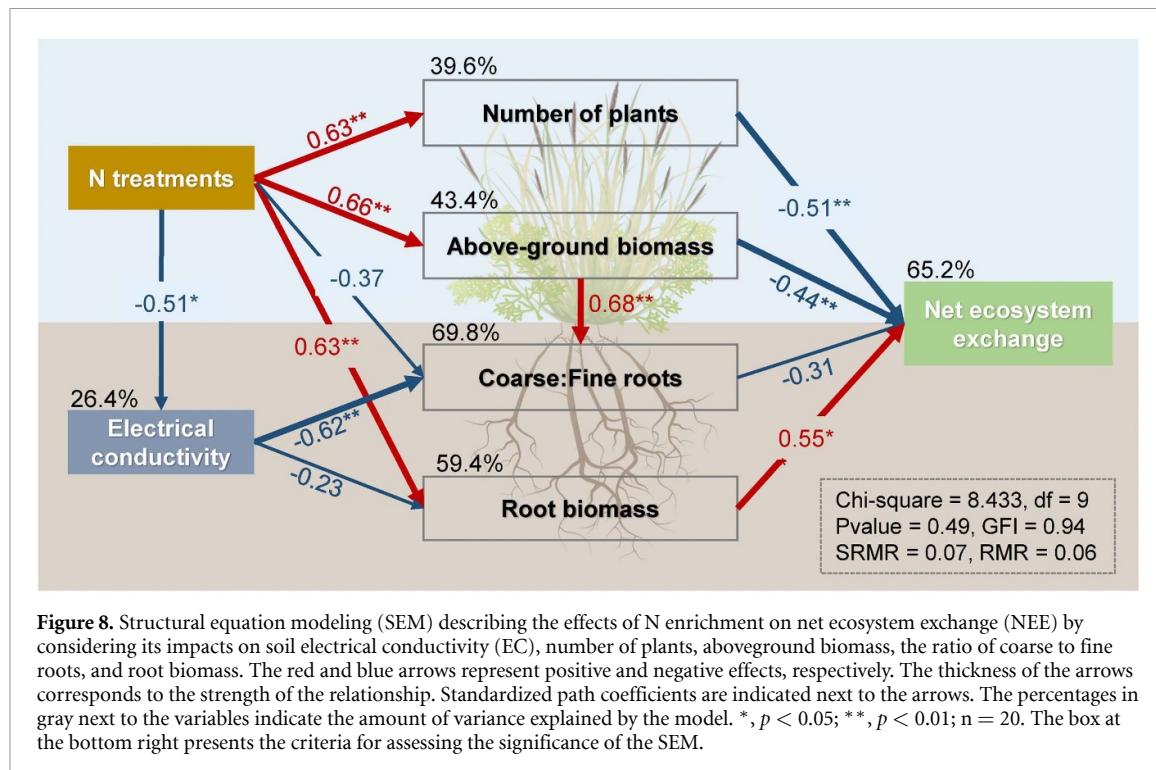


Figure 8. Structural equation modeling (SEM) describing the effects of N enrichment on net ecosystem exchange (NEE) by considering its impacts on soil electrical conductivity (EC), number of plants, aboveground biomass, the ratio of coarse to fine roots, and root biomass. The red and blue arrows represent positive and negative effects, respectively. The thickness of the arrows corresponds to the strength of the relationship. Standardized path coefficients are indicated next to the arrows. The percentages in gray next to the variables indicate the amount of variance explained by the model. *, $p < 0.05$; **, $p < 0.01$; $n = 20$. The box at the bottom right presents the criteria for assessing the significance of the SEM.

CO_2 fluxes. Third, N enrichment could modify CH_4 and N_2O fluxes, which are important components of the C cycle in wetlands. Related investigations on CH_4 and N_2O responses are still needed in the future to grasp the C-N interactions in coastal wetlands. Our study highlights the effects of N enrichment on the C cycling in a coastal wetland on the basis of long-term field experiments. We found level-dependent and positive impacts of N enrichment on the CO_2 sink in the coastal wetland, which is associated with N-induced shifts in soil and plants. Improving current C models by integrating the effects of N deposition on CO_2 sequestration is urgently and crucially needed. This improvement would contribute greatly to the global C budget and its accurate assessment.

5. Conclusions

Coastal wetlands have a crucial role in mitigating climate change as they serve as important C sinks. The impacts of N enrichment on ecosystem CO_2 cycling, however, still need to be identified, partly due to insufficient field experiments. We investigated the patterns and mechanisms of the effects of N on the sequestration of CO_2 in a YRD coastal wetland in China. We found that long-term N enrichment altered soil properties, plant growth, species composition, and patterns of root biomass. All these changes jointly and interactively stimulated the CO_2 fluxes in coastal wetlands. The N impacts depended on the level of N enrichment, indicating a nonlinear response of the functioning of the ecosystem CO_2 sink to nutrient stress in the wetland. Global

N deposition is projected to increase, so our study highlights the potential of stronger CO_2 sinks in coastal wetlands under scenarios of future climate change.

Data availability statement

The data that support the findings of this study are openly available at the following URL/DOI: <https://doi.org/10.5281/zenodo.11316363>.

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Author contributions

Baohua Xie and Xiaoyue Wang designed the study. Wendi Qu carried out the experiment, conducted all the analyses, constructed the figures and tables, and wrote the first draft. All authors contributed to the interpretation of the results and the text.

Conflict of interest

The authors declare no conflict of interest.

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