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Impact of ocean acidification on microzooplankton grazing dynamics

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This study examines the potential impacts of projected atmospheric carbon dioxide (pCO_2) levels reaching 800 ppm by the end of the century, focusing on ocean acidification effects on marine ecosystems in the coastal areas of Bohai. We investigated how acidification affects the grazing patterns of microzooplankton using dilution techniques and ecophysiological methods. Our findings indicate that acidic conditions shift the phytoplankton community structure, changing dominant species. Elevated CO₂ concentrations reduced grazing pressure on phytoplankton, with less steep declines in growth rates at 800 ppm CO₂ (spring: 2.43 d⁻¹ vs. 2.16 d⁻¹, summer: -0.46 d⁻¹ vs. -0.73 d⁻¹, autumn: -0.45 d⁻¹ vs. -0.90 d⁻¹) and significant decreases in grazing pressure percentages (%Pp from 0.84 to 0.58 and %Pi from 0.64 to 0.46). Short-term acid exposure significantly increased superoxide dismutase activity in both microplankton (from 0.03 to 0.08 U mg⁻¹, p<0.01) and nanoplankton (from 0.05 to 0.09 U mg⁻¹, p<0.001), indicating an adaptive response to oxidative stress. These results highlight that elevated CO2 levels primarily boost phytoplankton growth by reducing microzooplankton grazing pressure, resulting in higher growth rates and a shift towards smallersized phytoplankton, reflecting complex short-term ecological responses to acidification. Further research is needed to understand the long-term effects of ocean acidification on microzooplankton and their role in marine secondary productivity.

KEYWORDS

acidification, microzooplankton, grazing, superoxide dismutase activity, species composition

1 Introduction

Ocean acidification is a decrease in the pH value of the ocean surface caused by carbon dioxide emissions, which has a profound impact on global biogeochemical cycles (Gattuso et al., 2018). This process disrupts marine ecosystem biogeochemical cycles, impacting nutrient cycling, productivity, and altering the composition of food chains and webs (Flynn et al., 2012). Ocean acidification adversely affects the survival and reproductive success of

specific biological populations. As a result, these impacts extend to the broader ecological communities in which these populations exist (Pedersen and Hansen, 2003).

Advancements in acidification research have increasingly corroborated the adverse impacts of pH reduction on the survival, fecundity, and feeding behavior of microzooplankton, with implications for the oceanic carbon cycle and the transfer efficiency of energy to higher trophic levels (Hammill et al., 2018; Nelson et al., 2020). Nonetheless, investigations into the ecological ramifications of altered microzooplankton feeding behaviors under acidified conditions-particularly their protracted effects within natural environs-remain scant. Microzooplankton, pivotal in trophic cascades, bridge the microbial loop and the classical food chain (Azam et al., 1983; Caron and Hutchins, 2013). Prior inquiries have predominantly examined the influence of ocean acidification on the grazing rates of single microzooplankton species, with limited exploration into the impacts on natural communities feeding behaviors, thereby posing certain constraints. When organisms are exposed to high concentrations of CO₂, they may struggle to reduce the internal CO2 concentration. This leads to an accumulation of CO₂ and H⁺ in tissues and body fluids, which inhibits oxygen delivery. This condition is known as hyperacidemia. The physiological responses to hyperacidemia include impaired oxygen transport and delivery, which can significantly affect the organism's overall health and functionality (Pörtner, 2008). Particularly, physiological responses such as microzooplankton sensitivity to hypercapnia are elusive in natural assemblages. Meunier et al. (2017) delineated both the direct and indirect effects of ocean acidification on a single species. Directly, acidification may induce a decrement in zooplankton pH value and enzymatic activity, thereby impinging on physiological functions. For calcifying taxa, for instance, acidification could erode their calcium carbonate structures (Christopher E. et al., 2021). The study reveals that ocean acidification leads to an increase in reactive oxygen species (ROS) and malondialdehyde (MDA) levels, indicative of heightened oxidative stress in marine organisms. Notably, the activities of enzymes such as superoxide dismutase (SOD), catalase (CAT), and alkaline phosphatase (AKP) initially rise before decreasing, illustrating a complex, nuanced response to acidified conditions. This pattern highlights the potential threat posed by ocean acidification, not only affecting physiological functions but also compromising immune defense mechanisms in marine life. Furthermore, these physiological changes have broader ecological implications. As (Ramaekers et al., 2022) suggested, zooplankton growth may be indirectly affected by the diminished quality of primary producers in environments with elevated pCO_2 levels. This cascading effect from molecular changes to ecosystem dynamics underscores the far-reaching impact of ocean acidification. These observations suggest that the repercussions of ocean acidification on zooplankton are potentially twofold. This is linked to an associated decline in the quality of phytoplankton.

The response of the phytoplankton community to changes in environmental conditions may have the most significant indirect effect on the abundance, activity, and community composition of microzooplankton in the context of climate change. Phytoplankton serves as the primary food source for phytophagous microzooplankton, which directly or indirectly influences the quantity and composition of dissolved and particulate substances that enter the microbial food web, thereby serving as substrates for bacterial growth (Caron and Hutchins, 2013). Consequently, comprehending the impact of ocean acidification on the grazing rate and community dynamics of microzooplankton is essential for understanding the repercussions on both secondary productivity and primary productivity.

Contemporary research posits that ocean acidification may modify the feeding selectivity and efficiency of microzooplankton, subsequently affecting their ecological niche and functional role within the food web (Hammill et al., 2018). For instance, elevated acidification levels have been observed to skew certain microzooplankton preferences towards smaller prey, potentially restructuring community composition and trophic interactions (Hurd et al., 2018). Moreover, emerging research into the effects of acidification on the SOD activities in marine life, including microzooplankton, underscores the nuanced relationship between environmental stress factors and antioxidant defense mechanisms.

This study aims to fill the gap in current academic knowledge regarding the impact of ocean acidification on the feeding dynamics of microzooplankton communities. By employing a comprehensive methodology that merges controlled laboratory experiments with in-depth field studies, our aim is to accurately quantify how microzooplankton grazing rates vary under different acidification conditions. Moreover, this research will assess the broader implications of these changes on the structural and functional aspects of marine food webs. The expected outcomes of this study are poised to enhance our understanding of the ecological consequences of ocean acidification, providing crucial insights for the effective management and conservation of marine ecosystems.

2 Methods

2.1 Sample collection

Experiments were conducted at multiple points in time, including July, August, September, October of 2022, as well as March, April, and April of 2023. May and June. The sampling was collected surface seawater (<5m) from the coastal area of Bohai Sea in Tianjin, China (38°58'N, 117°50'E), filtered through a 200µm sieve, and filled in a 25L polyethylene bucket soaked in HCl in advance, which was quickly brought back to the laboratory for the preparation of acidified water. Instant temperature and pH were measured by S210-KpH meter (Mettller Toledo). For baseline community analysis, a 500ml aliquot of the initial water sample was reserved to examine the native microzooplankton and phytoplankton community structures. To analyze the fractionated chlorophyll a content, specific volumes of seawater were immediately filtered through 20µm and GF/F filter membrane (Whatman 47mm). The filter we use is Pall (4204), GFF membrane with a diameter of 25mm 0.2µm. Each fraction was then prepared in triplicate and stored at -20°C in a dark environment to prevent degradation.

2.2 Preparation of acidified seawater

Part of the collected seawater was filtered using a capsule filter and a GF/F membrane to make PFW (particle-free water). The PFW and initial seawater in the acidification group were mixed with high-pressure CO_2 and CO_2 enricher (CE100) for sufficient aeration to control the partial pressure of CO_2 at 800 ppm. The nonacidification group used a fish pump for full aeration (approximately 30 minutes), with a standard atmospheric pressure of 400 ppm. The instant pH was measured by S210-KpH meter (Mettller Toledo) calibrated with pH 7 and 10 buffer solutions during the bubbling process. When the water sample was stable and the pH did not change, the aeration ended. The water samples were maintained at a constant temperature during the entire process.

2.3 Microzooplankton grazing experiment

In this experiment, we employed the dilution method, as outlined by (Landry and Hassett, 1982), to assess the grazing rate of microzooplankton and the intrinsic growth rate of phytoplankton. The particle-free water (PFW) was carefully mixed with the initial seawater sample in varying proportions: 20%, 40%, 60%, 80%, and 100%. During this process, we used the siphon method for sub-packaging to prevent bubble formation, ensuring that each culture flask was filled completely without leaving any air gaps.

Each dilution proportion was replicated in three parallel samples, forming the treatment groups. The 1.5 L culture flasks were incubated in an incubator, which was gently agitated every 2 hours to maintain uniform conditions. Light conditions simulated natural environmental cycles with a 12-hour light and 12-hour dark period. After the 24-hour incubation period, each culture flask was gently agitated to ensure uniformity, and the pH of the samples was accurately measured. To assess phytoplankton growth, chlorophylla samples were collected from each flask. This involved the graded filtration of a specific volume of seawater through filter membranes with varying pore sizes to separate different particle fractions. The chlorophyll-a samples were immediately stored in a dark environment at -20° C to prevent degradation.

The chlorophyll *a* concentration, derived from these samples, served as a key indicator for estimating the apparent growth rate (AGR) of phytoplankton. This method allowed us to measure changes in phytoplankton biomass as a response to varying microzooplankton grazing pressures under the different dilution treatments. The careful preservation and analysis of these samples were critical for accurately determining the ecological dynamics between microzooplankton and phytoplankton in the context of our study. The growth of phytoplankton is expressed as formula:

$$P_t = P_o \,\mathrm{e}^{(\mu - dm)t} \tag{1}$$

where *t* is the duration of the incubation in days, P_o and P_t represent the initial and final concentrations of Chl a, respectively.

The apparent growth rate of phytoplankton

$$AGR (d^{-1}) = \frac{\ln(P_t/P_0)}{t}$$
$$= \mu - m$$
(2)

was calculated and linearly regressed with the actual dilution factor (ADF). The intercept of the equation was the intrinsic growth rate of phytoplankton (μ), and the slope was the grazing rate of zooplankton (m). According to the Equation 3, the net growth rate of phytoplankton (NGR), the grazing pressure of microzooplankton on phytoplankton standing stock (P_i), and the grazing pressure of microzooplankton on primary productivity (P_p) can be obtained:

$$NGR = \mu - m \tag{3}$$

$$P_i = 1 - e^{-m} \times 100 \,\% \tag{4}$$

$$P_p = \frac{e^{\mu} - e^{\mu - m}}{e^m - 1} \times 100\%$$
 (5)

2.4 Determination of SOD activity

The samples were immediately stored in a refrigerator (Haier DW-86L386) at -80° C after collection and filtration to maintain the activity before the determination of various indicators. SOD activity is a proxy for the cellular defense against reactive oxygen species, which are often elevated under stress conditions such as high CO₂. For the determination of SOD activity, we employed a commercially available SOD assay kit, utilizing the WST-1 method, sourced from the Nanjing Jiancheng Institute of Bioengineering. The WST-1 method is a widely recognized technique for its sensitivity and accuracy in detecting SOD activity. It involves the use of a watersoluble tetrazolium salt that produces a water-soluble formazan dye upon reduction with a superoxide anion, allowing for the quantification of SOD activity based on the rate of the reduction process.

For the measurements, we used GF/F filters with a pore size of 0.2 μ m and mesh screens with a pore size of 20 μ m, consistent with those used for chlorophyll-a measurements. The samples were incubated at 37°C for 20 minutes according to the kit's instructions, and absorbance was measured at 450 nm using a microplate reader. For SOD activity measurements, 5 ml samples were taken from each dilution gradient and averaged, excluding the initial seawater samples, focusing on comparing the differences between 400 ppm and 800 ppm CO₂ concentrations.

2.5 Analysis of microzooplankton and phytoplankton community

Microplankton water samples were fixed with formaldehyde (final concentration 1%). Phytoplankton and microzooplankton samples were analyzed under RX50 optical microscope. After each 500mL sample was settled for one week, the supernatant was removed and shaken well. On the premise of ensuring that the microzooplankton and phytoplankton in the water sample were not shaken and broken, the species identification and individual counting were performed at $200 \times$ and $400 \times$ magnification.

The Microplankton and phytoplankton species were identified according to Isamu Y (Isamu, 1959), and Sun (Jun et al., 2002).

Phytoplankton community diversity was calculated according to the Shannon–Wiener (S–W) diversity index (*H*') (Shannon, 1948):

$$H' = -\sum_{i=1}^{s} p_i \log_2 P_i$$
 (6)

where *Pi* represents the relative cell abundance of a species, *i* represents the numbers of the species, *S* represents the number of total species of phytoplankton in the samples collected.

The phytoplankton dominance index (*Y*) is calculated as follows:

$$Y = \frac{n_i}{N} f i \tag{7}$$

ni: the number of a single species; *N*: the number of all species; *fi*: the frequency of occurrence of the species in the sample.

2.6 Statistical analysis

Canonical Correspondence Analysis (CCA) was conducted using Canoco 5.0 to analyze the cell abundance of species and environmental factors, with data sets transformed using log10(x +1). Hierarchical clustering was applied to perform a cluster analysis of phytoplankton, with the dissimilarity between different phytoplankton groups calculated using Primer 6 (version 6.1.12.0).

All measurements were replicated at least three times, and the data were expressed as mean values \pm standard deviation (SD). ANOVA test was performed to compare the differences of various environmental and biological parameters between the 2 groups in present study. Significant differences were indicated at * p<0.05, ** p<0.02. Statistical analysis was performed using SPSS v. 16.0.

3 Result

3.1 Seasonal variation in phytoplankton community response to CO₂ concentrations

3.1.1 Impact of elevated CO₂ concentrations on phytoplankton size class distribution

At a CO_2 concentration of 400 ppm, micro-phytoplankton were observed to have chlorophyll a proportions ranging from 42.60% to 72.19%. In contrast, nano-phytoplankton exhibited proportions varying from 27.81% to 57.40%, as detailed in Figure 1. This variability highlights the dynamic nature of the phytoplankton community structure, characterized by significant fluctuations in the dominance between these two size classes.

In response to an increased CO_2 concentration of 800 ppm, a distinct shift in the phytoplankton community was noted. Micro-phytoplankton chlorophyll a proportions declined to a range of



28.40% to 58.01%, while those of nano-phytoplankton increased to between 41.99% and 71.60%. This alteration in the size class distribution underscores a trend towards greater dominance of nano-phytoplankton under higher CO_2 conditions (Figure 1). The observed shift suggests an ecological adaptation within the community, favoring smaller size classes as CO_2 concentrations rise.

3.1.2 Spring responses

During the spring (Figure 2), at the standard atmospheric level of 400 ppm CO₂, *Chaetoceros tortissimus* and *Chaetoceros lauderi* were amongst the most dominant, contributing to 14.87% and 13.29% of the community, respectively. At an elevated pCO_2 of 800 ppm, the pie chart illustrates an adjustment in relative abundance with *Chaetoceros pelagicus* and *Chaetoceros lorenzianus* becoming more pronounced, possessing dominance (Y) indices of 0.14 and 0.12 (Table 1), respectively. This quantitative shift suggests a change in the relative fitness of these species under higher CO₂, with potential implications for the structure and function of the spring phytoplankton community.

During the spring (Figure 3 A, S), under 400 ppm CO_2 conditions, phytoplankton abundance is around 10⁹ cells/L. However, under 800 ppm CO_2 conditions, the abundance drops to approximately 10⁸ cells/L. This suggests that elevated CO_2 levels may suppress phytoplankton growth during the early spring months.

3.1.3 Summer responses

The summer season displayed a broader change in community structure with increasing pCO_2 . While *Chaetoceros curvisetus* remained relatively dominant at both CO_2 levels (0.28 at 400 ppm and 0.28 at 800 ppm), the most noticeable change was the ascent of *Guinardia delicatula*, whose dominance index (*Y*) surged from 0.41 at 400 ppm to 0.50 at 800 ppm (Table 1). Figure 2 correspondingly reflect this ascendance, signaling a community



restructuring likely induced by the physiological or ecological preferences of *G. delicatula* under acidified conditions.

A substantial increase in phytoplankton abundance is observed in summer (Figure 3 M, J). The peak abundance reaches up to 10^{10} cells/L under 400 ppm conditions and as high as 10^{11} cells/L under 800 ppm conditions, indicating a positive response to elevated CO₂ levels and potentially enhancing phytoplankton productivity during this period of

peak growth. Phytoplankton abundance remains high, around 10^{10} cells/L for both $\rm CO_2$ conditions, suggesting that the growth-promoting effect of elevated $\rm CO_2$ persists throughout the summer.

3.1.4 Autumn responses

The autumn season marked a particularly interesting shift, with the dominance (Y) of *Akashiwo sanguinea* declining slightly from

Sample ID	400ppm	Dominance Index (Y)	800ppm	Dominance Index (Y)
Sp-1	Chaetoceros tortissimus	0.15	Chaetoceros pelagicus	0.13
	Chaetoceros lauderi	0.13	Chaetoceros lorenzianus	0.12
	Chaetoceros pelagicus	0.09	Chaetoceros paradoxus	0.12
	Chaetoceros lorenzianus	0.09	Chaetoceros decipiens	0.1
	Chaetoceros decipiens	0.08	Leptocylindrus danicus	0.09
Sp-2	Chaetoceros pelagicus	0.11	Chaetoceros paradoxus	0.14
	Chaetoceros tortissimus	0.11	Chaetoceros pelagicus	0.14
	Chaetoceros lauderi	0.11	Chaetoceros lorenzianus	0.12
	Chaetoceros lorenzianus	0.09	Chaetoceros decipiens	0.1
	Chaetoceros paradoxus	0.08	Chaetoceros lauderi	0.09
Su-1	Chaetoceros curvisetus	0.28	Chaetoceros diadema	0.28
	Chaetoceros diadema	0.16	Chaetoceros curvisetus	0.21
	Chaetoceros lorenzianus	0.09	Chaetoceros lorenzianus	0.21
	Chaetoceros debilis	0.09	Chaetoceros lauderi	0.06
	Chaetoceros tortissimus	0.08	Chaetoceros teres	0.06
Su-2	Guinardia delicatula	0.41	Guinardia delicatula	0.5
	Skeletonema costatum	0.17	Leptocylindrus danicus	0.16
	Leptocylindrus danicus	0.12	Nitzschia closterium	0.09
	Nitzschia.spp1	0.07	Skeletonema cf costatum	0.05
	Nitzschia closterium	0.06	Pseudonitzschia delicatissima	0.05
Au-1	Akashiwo sanguinea	0.77	Akashiwo sanguinea	0.7
	Gonyaulax bruunii	0.12	Gonyaulax bruunii	0.11
	Ceratium tripos	0.05	Pseudonitzschia delicatissima	0.07
	Gonyaulax fratercula	0.02	Ceratium tripos	0.04
	Gonyaulax polygramma	0.01	Ceratium furca	0.02
Au-2	Akashiwo sanguinea	0.42	Akashiwo sanguinea	0.4
	Chaetoceros tortissimus	0.08	Leptocylindrus danicus	0.13
	Ceratium furca	0.08	Ceratium furca	0.1
	Guinardia rotula	0.06	Thalassiosira nordenskioldi	0.06
	Skeletonema costatum	0.06	Pseudonitzschia pungens	0.05

TABLE 1 The top six dominant species and their dominance index in the three seasons (spring, summer and autumn) calculated using Equation 7.

0.77 at 400 ppm to 0.70 at 800 ppm, as detailed in Table 2. This change is visually represented in the pie charts (Figure 2). Despite maintaining a high dominance index, the community's composition exhibits a subtle yet discernible response to increased CO_2 levels. Potential shifts in nutrient uptake rates or grazing pressures are likely contributing factors to the observed succession.

In autumn (Figure 3 U, O), phytoplankton abundance remains high but starts to decline from summer peaks. The abundance is consistently around 10^{10} cells/L for both CO₂ levels, indicating a stabilization of growth. By late autumn, the abundance under 800 ppm CO₂ conditions is slightly higher than under 400 ppm, around 10^9 cells/L. This suggests a possible secondary growth phase or a delayed response to elevated CO_2 levels in the later part of the growing season.

3.1.5 Cross-seasonal growth responses

The cross-seasonal comparison showcases how elevated pCO_2 leads to differential responses within the phytoplankton assemblages, with some taxa like *C. pelagicus* showing an increase in dominance across multiple seasons (from 0.09 in spring at 400 ppm to 0.14 in spring at 800 ppm), whereas others, such as *G. delicatula*, demonstrate a pronounced seasonal response (surge in dominance index only during summer).



3.1.6 Species-specific growth responses

As depicted in Figure 4, it becomes evident that the response of phytoplankton species to elevated partial pressures of CO_2 is complex and varied. *C. pelagicus*, a species of interest, demonstrates a heterogeneous response to increased pCO_2 levels, with its growth rate fluctuating between minor increments (0.44), substantial increases (1.81), and instances of decline (-0.32). This variability suggests that the growth dynamics of *C. pelagicus* under elevated CO_2 conditions are influenced by a multitude of environmental or physiological factors, leading to oscillating growth patterns.

Similarly, *Chaetoceros paradoxus* exhibits significant growth rate enhancements, with a peak growth rate of 2.36 under elevated CO_2 concentrations. Nevertheless, this species also experiences minor negative growth rates (-0.06), indicating that its response to increased CO_2 levels is not consistently positive and may vary depending on specific environmental conditions or physiological states.

C. curvisetus reveals a pronounced variability in growth rates when subjected to enhanced CO_2 levels, with growth rates experiencing substantial escalations up to 3.474896, as well as notable reductions to -1.42. Such marked fluctuations underscore the possibility of divergent responses to varying environmental or physiological conditions induced by elevated CO_2 levels.

Leptocylindrus danicus displays enhanced growth under elevated CO_2 concentrations, achieving growth rates as high as 1.30, indicative of adaptability or resilience to these conditions. In contrast, *Gonyaulax bruunii* exhibits a consistent decline in growth

TABLE 2 Summary of microzooplankton grazing (md⁻¹) rates and phytoplankton growth (μ d⁻¹) for all the dilution experiments and relevant experimental parameters.

	m(d ⁻¹)	μ(d ⁻¹)	%Pp	%Pi	NGR
400ppm	1.05 ± 0.40	1.87 ± 0.40	0.75 ± 0.14	0.63 ± 0.13	0.82 ± 0.43
800ppm	0.57 ± 0.187	2.01 ± 0.44	0.50 ± 0.12	0.43 ± 0.11	1.44 ± 0.40
P	<0.05*	>0.05	<0.02**	<0.02**	<0.05*

* indicates p<0.05. ** indicates p<0.02.

rates across all conditions tested, with a significant reduction to -0.90, suggesting a detrimental response to increased CO₂ levels.

3.1.7 Physiological adaptations and environmental preferences

The analysis (Figure 5) identifies specific *Chaetoceros* species demonstrating varied responses to CO_2ParPr and pH levels. *Chaetoceros costatus* and *Chaetoceros decipiens* are associated with elevated pCO_2 , suggesting an adaptive preference or tolerance to high CO_2 conditions. Conversely, *C. curvisetus* and *C. lauderi* exhibit affinity for higher pH environments, indicating potential sensitivity to acidification, a byproduct of increased CO_2 levels. This delineation underscores the physiological heterogeneity within the phytoplankton community, with certain species potentially leveraging elevated CO_2 conditions to their advantage.

The CCA further reveals the thermal preferences of species like *Chaetoceros diadema* and *C. lorenzianus*, which align closely with the temperature vector, indicating a dependence on or adaptation to specific thermal conditions. The intermediate positioning of *Skeletonema costatum* and *L. danicus* between temperature and pH vectors suggests a broader ecological adaptability, allowing these taxa to thrive across a diverse range of environmental conditions.

Figure 5C accentuates the influence of pCO_2 on species distribution, particularly highlighting *Pseudonitzschia delicatissima* and *Peridinium pacificum* as taxa strongly affected by elevated CO₂ levels. Additionally, the proximity of *Ceratium furca* to the intersection of CO₂ concentration and temperature gradients points to the critical role of these combined environmental factors in determining species distribution patterns. This suggests that the interplay between CO₂ and temperature is pivotal in shaping the ecological dynamics of phytoplankton communities.

The CCA results collectively underscore the heterogeneity in physiological responses among phytoplankton to changes in pH and pCO_2 . This variability reflects the ecological diversity within the phytoplankton community, suggesting that certain taxa may possess adaptive advantages under scenarios of heightened CO_2 levels and subsequent acidification.



The dendrogram (Figure 6) exhibits a stratified pattern of community associations. Increasing the CO_2 concentration to 800 ppm, the clusters become more dispersed, signifying a divergent effect of acidification on the community structure. Notably, the 800 ppm autumn cluster demonstrates a broader spread, reflecting a potentially significant alteration of the phytoplankton assemblage in response to the combined effects of seasonality and elevated CO_2 levels. The spring samples (Figure 6), regardless of CO_2 treatment, show closer affinity, suggesting that the community composition during this period is less influenced by CO_2 variations and more by the inherent seasonal growth dynamics driven by factors such as increased light intensity and nutrient availability typical of spring blooms.

3.2 Microzooplankton grazing on phytoplankton in different pCO2 level

Throughout the seasons (Figure 7), a consistent trend emerges indicating that elevated CO_2 concentrations lead to a reduction in

grazing pressure on phytoplankton, as evidenced by the shallower slopes of apparent growth rate declines with increasing dilution factors. In spring, the decrease in growth rates was notably more pronounced at 800 ppm CO₂ (2.43 d^{-1}) compared to 400 ppm (2.16 d^{-1}), suggesting that higher CO₂ levels might mitigate grazing impacts, potentially due to acidification effects on zooplankton grazers or changes in phytoplankton palatability. This pattern persisted into summer and autumn, with reduced slopes observed at 800 ppm CO₂, indicating a sustained reduction in grazing pressure under elevated CO₂ conditions. Specifically, summer data ranged from -0.29 to -0.46 d⁻¹ at 800 ppm, compared to -0.66 to -0.73 d⁻¹ at 400 ppm, and in autumn, a less steep decrease was observed at 800 ppm ($-0.45 d^{-1}$) versus 400 ppm ($-0.90 d^{-1}$). These observations suggest that the interaction between phytoplankton growth and zooplankton grazing is significantly influenced by CO₂-nduced environmental changes, reflecting potentially complex shifts in aquatic food web dynamics. The seasonally varying response highlights the intricacy of ecological interactions in response to CO2-induced acidification, underscoring the importance of considering seasonal dynamics when assessing the ecological impacts of rising atmospheric CO₂ levels.

There was a clear decrease in %Pp and %Pi from 400 ppm to 800 ppm pCO_2 calculated using Equation 4 and Equation 5. At 400 ppm, the phytoplankton population faced significant grazing pressure (%Pp: 0.84 to 0.46; %Pi: 0.64 to 0.39), while at 800 ppm, the pressure was alleviated (%Pp: 0.58 to 0.30; %Pi: 0.462 to 0.25), suggesting a shift in the predator-prey dynamic likely influenced by ocean acidification (Figure 8).

The net growth rate (NGR), calculated using Equation 3 which indicates the overall growth of the phytoplankton population after accounting for losses due to grazing, was generally higher under elevated pCO_2 conditions (range: 0.99 to 1.88 d–1) compared to ambient CO_2 levels (range: 0.43 to 1.38 d⁻¹) (Table 2). To examine the potential CO_2 fertilization effect calculated using Equation 1 and Equation 2, we used the intrinsic growth rates (μ) for evaluation, as μ represents growth in the absence of grazing/mortality. Our findings show that μ did not shift significantly with treatment, suggesting that the changes in NGR are more likely related to variations in predation pressure rather than enhanced growth due to CO_2 fertilization.

The patterns discerned across different CO_2 treatments elucidate a significant dynamic where increased partial pressure of



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 CO_2 appears to mitigate grazing pressures on phytoplankton, facilitating enhanced net growth rates. This phenomenon is likely attributed to a decrease in the grazing efficiency of microzooplankton under higher CO_2 conditions. Consequently, these findings suggest a potential reduction in zooplankton's topdown control over phytoplankton populations, which could be the result of direct effects of elevated CO_2 on the physiological or behavioral aspects of both phytoplankton and zooplankton, as well as indirect effects on their interactions.

The correlation matrix analysis (Figure 9) reveals a nonuniform response of chlorophyll content to increased CO₂ levels. Nano-phytoplankton chlorophyll content (P-Chl-a) demonstrates significant correlations with the diversity of microzooplankton (MZP H') calculated using Equation 6, which could imply a linkage between primary producer biomass at the nano-scale and higher trophic levels. Micro-phytoplankton chlorophyll content (M-Chl-a) displays a less pronounced association with microzooplankton diversity. These observations suggest that the size-specific biomass of primary producers might influence microzooplankton community structures in disparate manners.

The significant negative correlation between pH and phytoplankton diversity (MPP H') across CO_2 treatments could be indicative of the direct impact of ocean acidification on phytoplankton community composition. It is apparent that as CO_2 concentration increases and pH decreases, phytoplankton diversity tends to increase, possibly due to differential sensitivities of species to pH changes, leading to shifts in community structure.

The Mantel's test results elucidate that the connections between environmental factors and the two CO_2 concentration levels are most pronounced for pH, highlighting acidification as a direct and significant consequence of elevated CO_2 . However, the direct effects of CO_2 on nano-phytoplankton and micro-phytoplankton chlorophyll contents appear to be less significant, suggesting that the chlorophyll concentration within these size classes may be regulated by a suite of indirect interactions and not solely by changes in CO_2 concentration.

3.3 Effect of acidification on microzooplankton superoxide dismutase

The activity of the antioxidant enzyme SOD in micro and nanoplankton was quantified to assess the oxidative stress response to different CO₂ concentrations (Figure 10). In microplankton, SOD activity increased significantly from 400 ppm to 800 ppm CO₂ concentrations, with values rising from an average of 0.03 U mg⁻¹ to 0.08 U mg⁻¹(p<0.01). The marked increase in SOD activity suggests that microplankton experience heightened oxidative stress at elevated CO₂ levels, triggering a pronounced cellular defense mechanism. For nanoplankton, a similar trend was observed. The SOD activity at 800 ppm CO₂ was substantially higher than at 400 ppm, with values averaging around 0.09 U mg⁻¹ compared to 0.05 U mg^{-1} at the lower CO₂ concentration (p<0.001), reinforcing the finding that nanoplankton also exhibits a strong antioxidative response to increased CO2. The observed increase in SOD activity in both micro and nanoplankton under elevated CO_2 conditions suggests that these communities may be actively countering the augmented oxidative stress associated with ocean acidification. This physiological response has implications for the resilience and



adaptive capacity of planktonic organisms in the face of ongoing environmental changes.

4 Discussion

4.1 Impact of ocean acidification on community composition

The availability of inorganic carbon significantly impacts phytoplankton photosynthesis and the activity of carbon concentrating mechanisms (CCMs). While a decrease in CCM activity can enhance the carbon fixation rate, conserving energy for cellular growth and photosynthesis, thus promoting phytoplankton growth (Thangaraj and Sun, 2021), our findings indicate that the observed increases in net growth rate (NGR) and total phytoplankton abundance are more likely attributable to reduced grazing pressure rather than a direct CO_2 fertilization effect. This conclusion is supported by the relatively stable intrinsic growth rates (μ) observed under varying CO_2 treatments. Although different algal populations exhibit varying preferences and tolerances towards carbon dioxide, which may drive future shifts in phytoplankton community composition in the ocean (Caron and Hutchins, 2013), the immediate impact on NGR appears to be predominantly influenced by changes in predation dynamics.

In previous studies, it was believed that pCO_2 had no discernible impact on the studied dinoflagellate and copepod species. However, in natural communities, some degree of physiological alteration in microzooplankton was observed (Meunier et al., 2017), aligning with the findings of Flynn et al. (2012). Even in cases where zooplankton is not directly affected by CO_2 aeration-induced



to Pp. * indicates p<0.05. ** indicates p<0.02.

disturbances, short-term exposure to high carbon dioxide levels can still induce Hyperacidemia. Rose et al. (2009) observed that shortterm increases in pCO_2 and temperature led to higher microzooplankton abundance and grazing rates, indicating a strong response to these variables in the short term. Suffrian et al. (2008) similarly found that short-term pCO_2 variations influenced microzooplankton grazing, although long-term effects were less pronounced.

Upon detailed examination of the predominant species across various seasons, as depicted in Figure 4, it becomes evident that the response of phytoplankton species to elevated partial pressures of CO_2 is complex and varied. The proximity of *Ceratium furca* to the intersection of CO_2 concentration and temperature gradients points to the critical role of these combined environmental factors in determining species distribution patterns. This suggests that the interplay between CO_2 and temperature is pivotal in shaping the ecological dynamics of phytoplankton communities. Additionally, the repercussions of ocean acidification on primary productivity are propagated through the food chain, indirectly influencing microzooplankton.

The experimental findings of this study demonstrate that when coastal surface seawater, rich in nutrients, is exposed to acidification





stress, there is a significant reduction in the proportion of microlevel phytoplankton community structure. Previous research by Collins (Collins et al., 2014) suggested that the growth rate of diatoms and eukaryotes in phytoplankton increased under acidification conditions. However, due to differential growth rates or selective grazing by microzooplankton, or acidification-induced reduction in silicon content of diatoms (Sharma et al., 2022), the grazing rate of microzooplankton decreased.

Furthermore, some scholars argue that algae with smaller particle sizes, such as ultramicro eukaryotic algae, Synechococcus, and Prochlorococcus, are more sensitive to elevated carbon dioxide levels (Hutchins et al., 2007; Levitan et al., 2010). Conversely, other studies propose that phytoplankton with larger particle sizes exhibit faster growth rates under acidification conditions, attributed to their smaller surface area ratio and reduced CO₂ diffusion (Sun et al., 2022). Nano-phytoplankton is more suitable for grazing by microzooplankton in size (Sun et al., 2003). However, due to the decrease in food level or the unfavorable state of zooplankton, the grazing rate still showed a downward trend. Based on the study by Mayers et al. (2020), both picophytoplankton and nanophytoplankton can be grazed by microzooplankton at similar rates. The research utilized flow cytometry to compare grazing rates on different phytoplankton groups, including picoeukaryotes, nanoeukaryotes, and coccolithophores. The findings indicate that the grazing rates were similar across these groups, demonstrating that microzooplankton do not exhibit significant preference or avoidance based on the size or calcification of the phytoplankton. This suggests that the possession of coccoliths by coccolithophores, such as Emiliania huxleyi, does not provide effective protection against microzooplankton grazing, challenging the hypothesis that calcification serves as a defensive mechanism against predation.

4.2 Stress response of microzooplankton in high CO₂ environment

When organisms are exposed to high levels of carbon dioxide, most organisms likely find it difficult to reduce the concentration of CO₂ in their bodies, leading to the accumulation of CO₂ and H⁺ in tissues and body fluids, thereby inhibiting the transport of oxygen (Pörtner, 2008), a physiological reaction known as Hyperacidemia. The SOD enzyme serves as an indicator of the oxidative stress response in organisms exposed to stressful environments. It can catalyze the disproportionation reaction of superoxide radicals to produce oxygen and hydroxyl radicals, thereby eliminating harmful superoxide radicals. In many organisms, the level of SOD is one of the important indicators to maintain the redox balance, which can characterize the effect of acidification environment on the growth of plankton to a certain extent. Some studies have shown that ocean acidification can reduce the activity of SOD in microzooplankton, thus affecting its antioxidant capacity. Under these conditions, the organism will produce ROS when threatened by external threats, and at the same time, it will also increase the activity of SOD to remove free radicals produced in the body to protect cells from oxidative damage (Wolfe-Simon et al., 2005).

In a study, the acidification conditions of seawater were simulated in the laboratory, and the activity of SOD enzyme in microzooplankton was measured. The activity of SOD in Prorocentrum dentatum and Karenia mikimotoi showed the same trend in the acidification culture system, that is, it increased first and decreased after two days (Zhang et al., 2022). In this study, the incubation time was only one day. It can be observed that the SOD activity of the mixed system of zooplankton and phytoplankton with two particle sizes increased significantly. The reason for this situation may be that the enzyme activity will cause its own adaptive regulation when it is oppressed by the living environment (Jahnke and White, 2003), so the SOD activity increases to eliminate the production of ROS. However, persistent damage may also lead to cell death and reduce the overall enzyme activity (Gao et al., 2019). Therefore, after a period of culture, the effect of stress on organisms is manifested as the inhibition of biological enzyme activity. Some studies also found that elevated CO₂ concentration inhibited the activity of SOD in copepods, resulting in oxidative stress in copepods (Zhang et al., 2016). Other studies Meunier et al. (2017) also believe that ocean acidification may destroy the redox balance of microzooplankton (Melzner et al., 2009), making it more vulnerable to oxidative stress. Another study also discussed the effect of ocean acidification on the SOD level of microzooplankton. This study suggested that ocean acidification may have a negative impact on the health and survival of microzooplankton, because the decrease of SOD activity may lead to a decrease of antioxidant capacity, thus increasing the risk of cell damage (Zhang et al., 2012). Of course, in addition to SOD activity, ocean acidification has other effects on the reduction of grazing rate of microzooplankton, such as their metabolic rate and growth rate, but this effect is rapid and limited (Dam et al., 2021), and the adaptation mechanism of microzooplankton to ocean acidification environment needs further study.

From the physiological and biochemical point of view, ocean acidification may lead to a decrease in the level of SOD in plankton in the short term, thus affecting the antioxidant capacity of microzooplankton. In addition, ocean acidification may also have a negative impact on the grazing rate of microzooplankton, which may further affect the survival and ecosystem function of microzooplankton. To understand these effects, it is necessary to further study the response mechanisms under different species and environmental conditions.

4.3 The long-term impact of ocean acidification on the survival and ecological functions of microzooplankton

Figure 11 visually summarizes our findings, illustrating the complex ecological responses in the marine environment to anthropogenic CO_2 emissions. Key trends include an increase in phytoplankton growth, primarily driven by reduced grazing pressure rather than the increased availability of carbon for photosynthesis. This reduction in grazing pressure allows for higher residual growth rates of phytoplankton. The growth is associated with shifts in community structure, notably a decrease in community particle size, suggesting a shift towards smaller-sized phytoplankton as a direct or indirect consequence of acidified conditions. The diagram also indicates altered grazing rates of microzooplankton, represented by the arrow, potentially reflecting an adaptive response to changes in prey availability or composition.

In the study by Aberle et al. (2013), while long-term pCO_2 changes did not significantly affect the composition and diversity of the microzooplankton community, short-term pCO_2 variations elicited different ecological responses. The study highlighted that

under high pCO_2 conditions, there were observable shifts in phytoplankton community structure, such as the positive effect on dinoflagellates. However, these changes were often transient and did not persist over longer periods, possibly due to the high buffering capacity of complex natural systems and the inherent variability in phytoplankton species composition.

In the short term, pCO_2 -induced alterations can rapidly affect phytoplankton population dynamics, thereby influencing the availability of food for microzooplankton and their grazing behaviors. These short-term ecological shifts, although not sustained in the long term, indicate the microzooplankton community's ability to quickly respond to environmental stressors. This suggests that while microzooplankton exhibit high tolerance to long-term ocean acidification, short-term changes in pCO_2 levels can significantly impact their community dynamics.

This affects the productivity and stability of microzooplankton and phytoplankton in the acidified environment of seawater. As the grazing rate of microzooplankton decreased, the growth rate of phytoplankton increased, and as shown in Table 2, $m < \mu$, the apparent growth rate of phytoplankton exceeded the grazing rate of microzooplankton, so the grazing pressure of microzooplankton on the standing stock of phytoplankton decreased. At the same time, the grazing pressure of microzooplankton on the potential productivity of phytoplankton is reduced, which means that in the acidified environment of seawater, the growth and production of phytoplankton will not be inhibited, which is related to the increase of net growth rate of phytoplankton, and the control effect of microzooplankton on phytoplankton is decreasing. Theoretical studies have shown that environmental stress has a strong impact on predators and a weak impact on prey, which is conducive to community rescue; on the contrary, it is not conducive to community rescue (Gilbert et al., 2014).



In the Mesocosm experiment designed by (Bell et al., 2019), it was observed that when the adaptability of zooplankton to water acidification was low, the predation stress on phytoplankton decreased, which may promote the evolutionary rescue of phytoplankton. However, if the predator was too sensitive, only phytoplankton survived, which may lead to the imbalance of energy flow and material cycle after a longer period.

5 Conclusion

In summary, the effects of ocean acidification on the grazing of microzooplankton are multifaceted. In this study, these effects led to a decrease in the standing grazing pressure of microzooplankton on phytoplankton and the grazing pressure of potential primary productivity. In the future studies, the effects of ocean acidification on secondary productivity and trophic communities still need more research and discussion.

Data availability statement

The raw data supporting the conclusions of this article will be made available by the authors, without undue reservation.

Author contributions

WS: Conceptualization, Formal analysis, Investigation, Methodology, Visualization, Writing – original draft, Writing – review & editing. XF: Software, Writing – review & editing. YH: Supervision, Writing – review & editing. JQ: Validation, Writing – review & editing. JS: Conceptualization, Methodology, Project

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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