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EDITED BY  
Zhangxi Hu,  
Guangdong Ocean University, China

REVIEWED BY  
Dajun Qiu,  
South China Sea Institute of  
Oceanology, Chinese Academy of  
Sciences, China  
Andreas Seger,  
University of Tasmania, Australia

\*CORRESPONDENCE  
Zhiming Yu  
zyu@qdio.ac.cn

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# Application of modified clay in intensive mariculture pond: Impacts on nutrients and phytoplankton

Lianbao Chi<sup>1,2,3</sup>, Yu Ding<sup>1,2,3,4</sup>, Liyan He<sup>1,2,3</sup>, Zaixing Wu<sup>1,2,3</sup>,  
Yongquan Yuan<sup>1,2,3</sup>, Xihua Cao<sup>1,2,3</sup>, Xiuxian Song<sup>1,2,3,4</sup>  
and Zhiming Yu<sup>1,2,3,4\*</sup>

<sup>1</sup>CAS Key Laboratory of Marine Ecology and Environmental Sciences, Institute of Oceanology, Chinese Academy of Sciences, Qingdao, China, <sup>2</sup>Laboratory for Marine Ecology and Environmental Science, Qingdao National Laboratory for Marine Science and Technology, Qingdao, China, <sup>3</sup>Center for Ocean Mega-Science, Chinese Academy of Sciences, Qingdao, China, <sup>4</sup>University of Chinese Academy of Sciences, Beijing, China

Nutrients and phytoplankton associated with mariculture development are important concerns globally, as they can significantly impact water quality and aquaculture yield. Currently, there is still insufficient information regarding the variations in nutrients and phytoplankton community of intensive mariculture systems, and effective treatment is lacking. Here, based on consecutive daily monitoring of two *Litopenaeus vannamei* ponds from July to October, the dynamic variations in nutrients and phytoplankton were elucidated. In addition, modified clay (MC) method was adopted to regulate the nutrients and phytoplankton community. The temporal variations in organic and inorganic nutrients presented fluctuating upward trends. Notably, organic nutrients were the dominant species, with average proportions of TON/P in TN/P were as high as 75.29% and 87.36%, respectively. Furthermore, a marked increase in the ratios of dinoflagellates to diatoms abundance were also observed in the control pond, concurrently with dominant organic nutrients, ascending N/P ratio and decreasing Si/N and Si/P ratios. In the MC-regulated pond, MC reduced the contents of both organic and inorganic nutrients. Furthermore, a distinct change pattern of dominant phytoplankton community occurred, with green algae becoming the most abundant phytoplankton in the MC-regulated pond. This study can provide new insights into an effective treatment for managing water quality and maintaining sustainable mariculture development.

## KEYWORDS

mariculture, dynamic variations, nutrient composition and structure, phytoplankton, modified clay

## Highlights

- Organic N and P were the dominant species of nutrients in mariculture shrimp ponds. Higher N/Si and N/P ratios were prominent over the production cycle.
- Variations in nutrients favored the potential predominance of dinoflagellates.
- Modified Clay effectively reduced nutrient contents and regulated the phytoplankton community.

## Introduction

Mariculture is one of the fastest-expanding sectors worldwide (Campbell and Pauly, 2013; Froehlich et al., 2018; Meng and Feagin, 2019), with global production reaching 87.5 million tons in 2020 (FAO (Food and Agriculture Organization of the United Nations), 2022). Notably, China has played a major role in this growth (Li et al., 2017). In China, pond farming is a dominant aquaculture farming system. For instance, the total production of pond farming continued to increase in 2019, contributing approximately 50% of national aquaculture production (Ministry of Agriculture and Rural Affairs of the People's Republic of China, 2020). However, the rapid development of mariculture has been accompanied by environmental pollution, such as nutrient and organic matter overload, harmful algal blooms, and drug residues, which pose threats to mariculture development (Briggs and Funge-Smith, 1994; Couch, 1998; Bouwman et al., 2013; Han et al., 2021). Currently, mariculture has become a significant and expanding cause of coastal nutrient enrichment and has influenced marine ecosystems of the receiving coastal waters (Bouwman et al., 2013; Li et al., 2017; Yang et al., 2017).

As is the case with all intensive farming systems, large quantities of excess feed and faeces increase the nutrient loading to water bodies, which cause the deterioration of water quality and restrict the sustainable mariculture (Alonso-Rodríguez and Páez-Osuna, 2003; Casé et al., 2008; Yang et al., 2017; Díaz et al., 2019). Notably, the accumulation of ammonia and nitrite will exert toxic effects on aquaculture organisms and cause disease proliferation in culture ponds (Casé et al., 2008; Glibert, 2016). Moreover, the excessive accumulation and unbalanced proportion of nutrients will influence the phytoplankton community and can cause the occurrence of 'harmful algal blooms (HABs)' (Glibert, 2016; Díaz et al., 2019). Increasing occurrences of HABs, such as *Karenia* spp., *dinoflagellates*, and *Aureococcus anophagefferens*, are leading to growing deleterious impacts including the poisoning, asphyxiation and even the death of mariculture organisms and human poisoning (Davidson et al., 2009; Gobler et al., 2011; Anderson, 2012; Brown et al., 2020). On the other hand, the

unbalanced proportion of nutrients can cause a significantly higher toxicity activity of phytoplankton (Johansson and Granéli, 1999; Hagström and Granéli, 2005). Furthermore, the discharge of mariculture effluents generates diverse effects on coastal waters (Bouwman et al., 2013). For instance, over 26% of the excess nitrogen in China's waters is likely a result of shrimp production alone (Meng and Feagin, 2019).

Modified clay (MC), produced from natural clays *via* the surface modification by inorganic or organic compounds, can effectively control HABs (Yu et al., 2017; Song et al., 2021). Currently, MC technology has been included as a national standard method to control HABs and is widely employed in China (Yu et al., 2017). Previous studies found that MC can not only remove algal cells but also reduce nutrients and organic matter contents, improve water quality and reduce the degree of eutrophication (Gao et al., 2007; Lu et al., 2017; Yu et al., 2017; Song et al., 2021). In addition, MC has no adverse effects on the survival and growth of typical economically marine organisms when used at appropriate dosages (Zhang et al., 2019; Song et al., 2021). Furthermore, in mitigating the blooms of toxin-producing dinoflagellates, MC can quickly reduce algal toxins in water (Hagström and Granéli, 2005; Lu et al., 2017; Li et al., 2019).

The variations in nutrients and phytoplankton associated with mariculture development are important concerns globally, as they can have a variety of potential impacts on the mariculture yield and the environment of receiving water. Currently, there is still insufficient information on the dynamic variations in nutrients and phytoplankton, and effective regulation treatment is lacking. In the present study, based on daily monitoring of nutrients and phytoplankton of intensive mariculture shrimp ponds in Laizhou Bay, the specific objectives were to: (i) elucidate the dynamic variations in nutrients and explore the potential responses of phytoplankton community to the variations in content, composition and stoichiometric ratio of nutrients, (ii) compare the variations in nutrients and phytoplankton community between an MC-regulated and control ponds and assess the regulation effects of MC. The results of this study are crucial for the comprehensive understanding of the variation patterns of nutrients and phytoplankton in the mariculture systems, and provide new insights into an effective regulation treatment for managing water quality and maintaining sustainable mariculture development.

## Materials and methods

### Study area

Intensive rearing *Litopenaeus vannamei* ponds were located in Dongying, Laizhou bay, China (118°55'E, 37°27'N), and two shrimp ponds were selected as the control pond and the MC-regulated pond, respectively (Figure 1). Each pond covered an

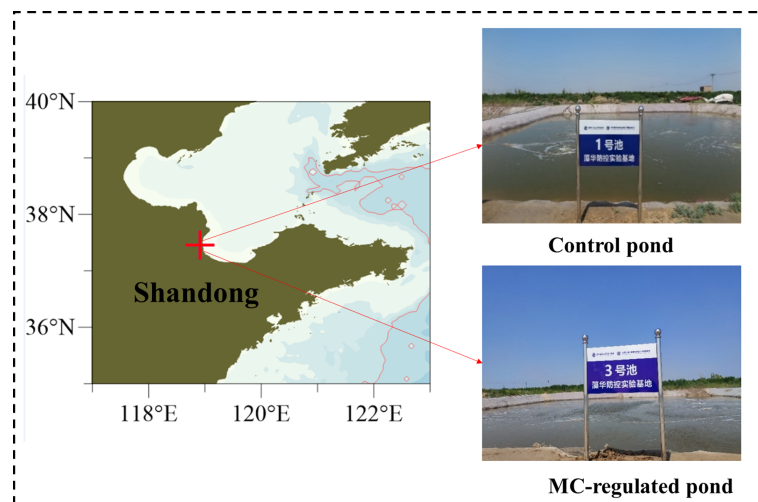


FIGURE 1  
The locations of mariculture shrimp ponds in the Laizhou bay.

area of approximately 1000 m<sup>2</sup>, with an average water depth of 1.5 m (shrimp stocking density:  $5 \times 10^4$  shrimp per pond, the average body length of shrimp: 1.6 cm). There were no water changes during the culture process and each pond was equipped with an aerator to ensure a suitable dissolved oxygen concentration. The control pond adopted the traditional culture method for breeding, while the MC-regulated pond adopted MC to regulate water quality and phytoplankton community. The elemental composition and content of the MC were determined by XRF, as shown in Table S1. The MC is prepared into a suspension with seawater and then sprayed on the pond in a boat with spraying equipment. And the spraying of the MC were determined according to the variations in water quality and phytoplankton community (Table S2).

## Sampling and analysis

Hydrographic parameters, including temperature and salinity, were measured by a YSI multiparameter water quality meter (YSI Ltd., USA). The water samples analyzed for nutrients, Chl *a* and 18S rDNA were collected at a depth of 0.5 m below the surface from multiple points (four corners and the center) in the pond every two days. In addition, parallel samples were collected every week. Samples for measuring total nitrogen (TN) and total phosphorus (TP) were collected and stored at -20°C in a sulfuric acid solution (50% v/v) with a final concentration of 0.2%. Water samples for the determination of dissolved inorganic nutrients were filtered under dim light through Whatman GF/F filter membranes (pore size: 0.68 μm). The filtrate was poured into 60 mL polyethylene bottles added to approximately 0.1 mL of chloroform and stored at -20°C. For the Chl *a* measurement,

100 mL of water sample was filtered using What man GF/F filters after initial filtration through a 200 μm nylon sieve, and the filter membrane was wrapped in stored in a dark environment at -20°C. Water samples for 18S rDNA were filtered through a 200 μm nylon sieve to eliminate the interference of zooplankton. Following filtration, the samples were passed through a 0.22 μm cellulose acetate membrane filter. The filters were collected in a centrifuge tube and stored in liquid nitrogen at -196°C.

In the laboratory, the concentrations of TN, TP, and dissolved inorganic nitrogen [DIN, including nitrate (NO<sub>3</sub><sup>-</sup>), nitrite (NO<sub>2</sub><sup>-</sup>), TAN (non-ionic ammonia (NH<sub>3</sub>) and ammonium ion (NH<sub>4</sub><sup>+</sup>)), dissolved inorganic phosphorous (DIP) and dissolved silicate (DSi) were measured with a SKALAR Flow Analyser (Skalar Ltd., Netherlands). The TON and TOP contents were calculated by the differential subtraction method, i.e. TON=TN-DIN, TOP=TN-DIP. Chl *a* was extracted in acetone, and its concentrations were determined by using a Trilogy fluorometer (Turner Design Ltd., USA).

## DNA extraction, PCR, and gene sequencing

DNA was extracted using the HiPure Soil DNA Kits (Magen, Guangzhou, China) according to the manufacturer's protocol. For 18S rDNA genes, universal primers 528F (5'-GCGGTAATTCCAGCTCCAA-3') and 706R (5'-AATCCRA GAATTCACCTCT-3') targeting the V4 regions were used for PCR (95°C for 2 min, followed by 35 cycles at 95°C for 30 s, 60°C for 45 s, 72°C for 90 s, and a final extension at 72°C for 10 min). Ampicons were extracted using 2% agarose gels and purified using

the AxyPrep DNA Gel Extraction Kit (Axygen Biosciences, Union City, CA, USA) according to the manufacturer's instructions. Purified amplicons were pooled in equimolar amounts and paired-end sequenced (PE250) on an Illumina platform according to the standard protocols.

## Data processing and analysis

The raw reads were further filtered using FASTP to obtain high-quality clean reads. Paired-end clean reads were merged as raw tags using FLASH, with a minimum overlap of 10 bp and a mismatch error rate of 2%. The clean tags subjected to specific filtering conditions were clustered into operational taxonomic units (OTUs) of  $\geq 97\%$  similarity using the UPARSE pipeline. All chimeric tags were removed using the UCHIME algorithm; finally, effective tags were obtained for further analysis. The tag sequence with the highest abundance was selected as the representative sequence within each cluster. The representative OTU sequences were classified into organisms by a naïve Bayesian model using the RDP classifier based on the SILVA database, with a confidence threshold value of 0.8. To further elucidate the abundance and richness of eukaryotic phytoplankton from the 18S rDNA results, the nonalgal OTUs, including Metazoa, Fungi, Apicomplexa, Intramacronucleata, Streptophyta, Postciliodesmatophora, Opalozoa, and unclassified data were removed. In addition, the relative abundance of each taxon was calculated by dividing the sequence number of OTUs of this group by the total sequence number of algae.

The ranking analysis of phytoplankton and environmental factors was performed using CANOCO for Windows 5.0 software package. To satisfy the assumptions of normality and homogeneity of variance, all species data and environmental data were subjected to  $\log(x + 1)$  transformation prior to multivariate analysis (Lepš and Šmilauer, 2003). According to the results of detrended correspondence analysis (DCA), redundancy analysis (RDA) was selected to determine the relationship between phytoplankton and environmental factors. The significance of the RDA ranking model was verified by the Monte Carlo permutation test.

## Results

### Temporal variations in nutrients

The TN and TP concentrations in the control and MC-regulated ponds varied in the ranges of 95.10–588.76  $\mu\text{mol/L}$ , 100.86–557.41  $\mu\text{mol/L}$  and 1.73–22.47  $\mu\text{mol/L}$ , 2.24–19.32  $\mu\text{mol/L}$ , respectively. Notably, the contents of TN (Figure 2A) and TP (Figure 2B) in both ponds exhibited a rapid and fluctuating upward trend over time. Overall, during the main culture period, the TN and TP concentration of the MC-regulated pond were

lower than those of the control pond. As for TON and TOP, the ranges in the control and MC-regulated ponds were 88.66–478.81  $\mu\text{mol/L}$ , 43.81–387.82  $\mu\text{mol/L}$ , and 0.81–21.42  $\mu\text{mol/L}$ , 1.85–18.60  $\mu\text{mol/L}$ , respectively. The mean concentrations of TON (Figure 2C) and TOP (Figure 2D) were significantly higher than those of DIN and DIP, which indicated that organic nutrients were the predominant species of nutrients in the water column of shrimp ponds. Similarly, significant temporal variations in TON and TOP concentrations were observed over the study period, which showed an increasing trend over time. Overall, in the late stage, the TOP concentration of the MC-regulated pond was lower than that of the control pond. The concentrations of DIN and DIP in the control and MC-regulated ponds varied in the ranges of 0.91–281.82  $\mu\text{mol/L}$ , 0.79–248.37  $\mu\text{mol/L}$ , and 0.22–5.62  $\mu\text{mol/L}$ , 0.19–3.75  $\mu\text{mol/L}$ , respectively. Significant temporal variations in the DIN (Figure 2E) and DIP (Figure 2F) were observed, with a fluctuating upward trend observed in the middle and late stages. Comparatively, the concentrations of DIN, especially DIP, in the MC-regulated pond were lower than those in the control pond during the study period. The DSi concentrations remained relatively stable during the study period (Figure 2G), and DSi concentrations in the control and MC-regulated ponds varied in the ranges of 1.55–8.96  $\mu\text{mol/L}$  and 1.81–8.86  $\mu\text{mol/L}$ .

The concentrations of TAN,  $\text{NO}_2^-$  and  $\text{NO}_3^-$  ranged as follows: 0.78–301.93  $\mu\text{mol/L}$ , 0.01–9.41  $\mu\text{mol/L}$  and 0.00–8.63  $\mu\text{mol/L}$ , respectively, in the control pond; while 0.70–239.71  $\mu\text{mol/L}$ , 0.00–6.23  $\mu\text{mol/L}$ , and 0.00–9.21  $\mu\text{mol/L}$ , respectively, in the MC-regulated pond. Notably, TAN was the predominant species of DIN, with significantly higher concentrations than those of  $\text{NO}_2^-$  and  $\text{NO}_3^-$ , and its average content accounted for 86.72% and 88.30% of the DIN concentration in the control and MC-regulated ponds, respectively (Figure 3A). Temporal variations in the TAN (Figure 3B) and  $\text{NO}_2^-$  (Figure 3C) concentrations of both ponds were observed, which exhibited an overall increasing trend during the study period. In addition, the TAN and  $\text{NO}_2^-$  concentrations were lower in the MC-regulated pond than in the control pond during the main study period. Differently, the concentrations of  $\text{NO}_3^-$  exhibited a different temporal variation (Figure 3D), decreasing in the early stage and then increasing in volatility during the middle and late stages.

### Temporal variations in Chl *a* concentrations and phytoplankton community

The Chl *a* concentrations ranged between 12.57–635.40  $\mu\text{g/L}$  and 12.25–489.93  $\mu\text{g/L}$  in the control and MC-regulated ponds, respectively. There were significant temporal variations in Chl *a* concentrations in the two ponds, with considerably lower concentrations observed during the initial stage than the other

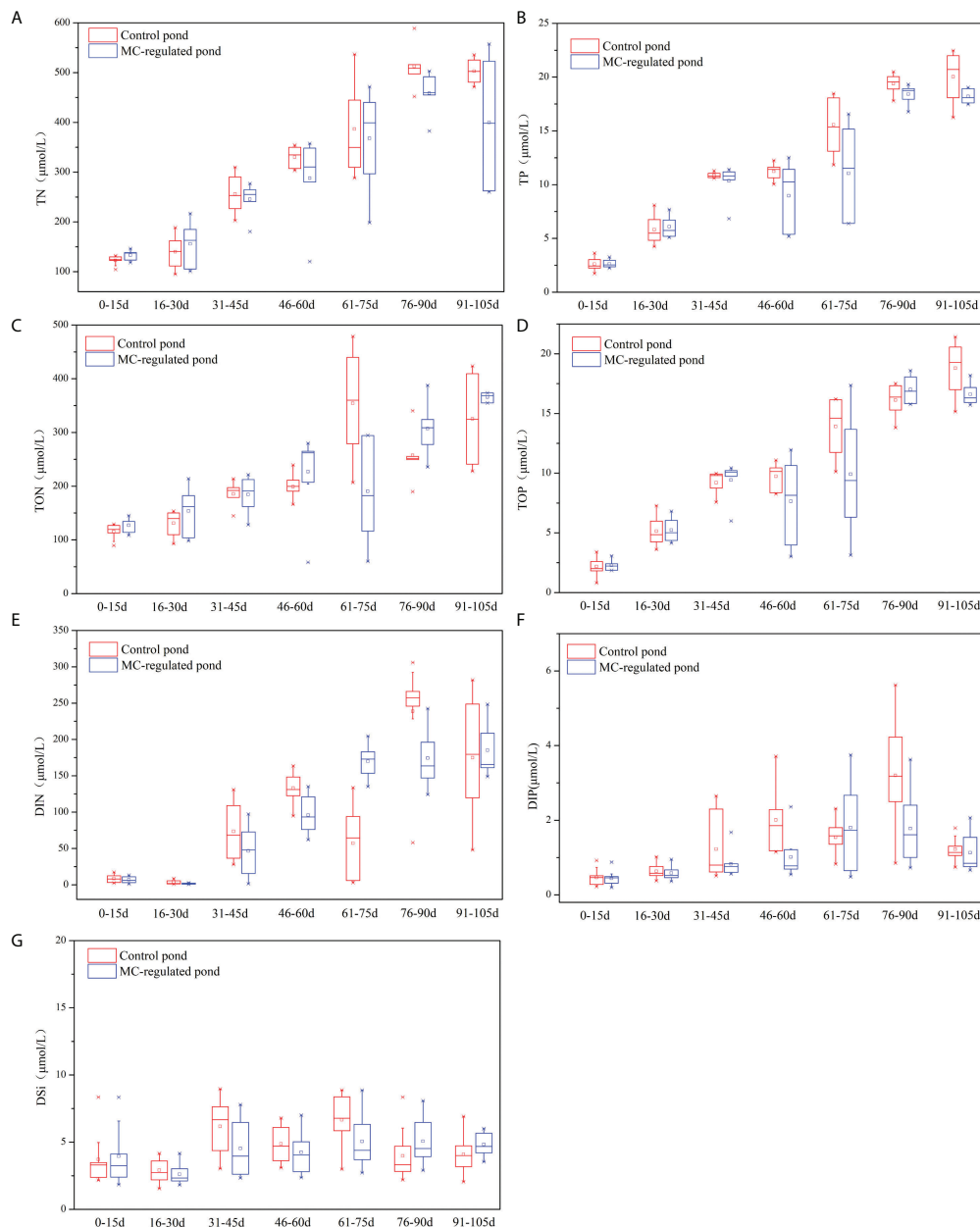
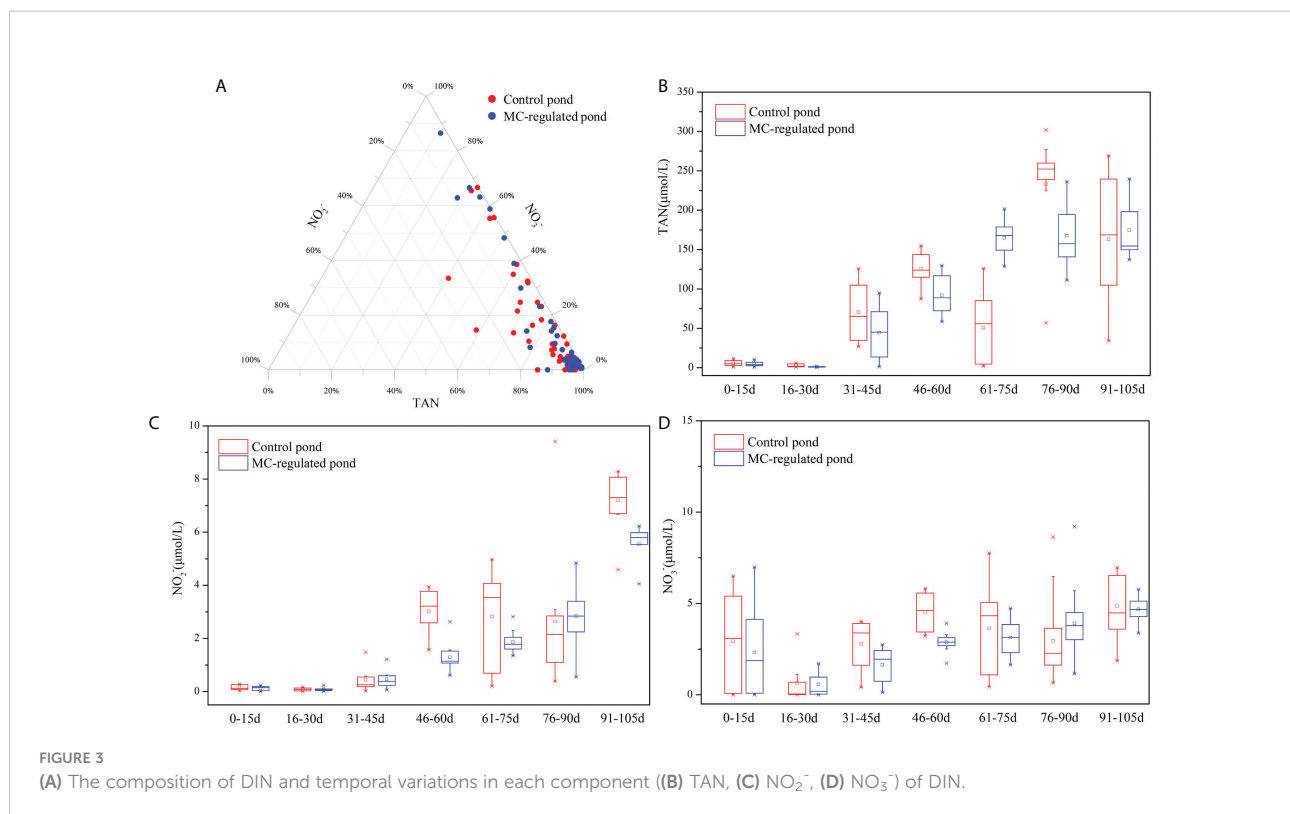


FIGURE 2  
Temporal variations in (A) TN, (B) TP, (C) DON, (D) DOP, (E) DIN, (F) DIP and (G) DSI concentrations in the water column.

two stages (Figure 4A). Comparatively, the variation of Chl *a* in the control pond exhibited greater fluctuations than that in the MC-regulated pond. As for phytoplankton community, a total of 2,728,684 effective tags by 18S rDNA gene sequencing were obtained, after quality screening, with an average N50 of 343 bp. After clustering based on a 97% similarity threshold and removing non-algal OTUs, 212 and 222 algal OTUs were obtained from the control and the MC-regulated ponds, respectively. Overall, six phyla, 16 classes, 22 orders, 29

families, 33 genera and 38 species were identified after annotation. Among them, the top ten algae with the highest abundance at the family level were defined as the dominant phytoplankton genera, and the abundance of families in the two ponds is shown in Figures 4B, C. In the initial stage, the phytoplankton community in both the control and the MC-regulated ponds were similar and dominated by diatoms, mainly including Stephanodiscaceae at the family level. Thereafter, the phytoplankton composition exhibited temporal variations, with

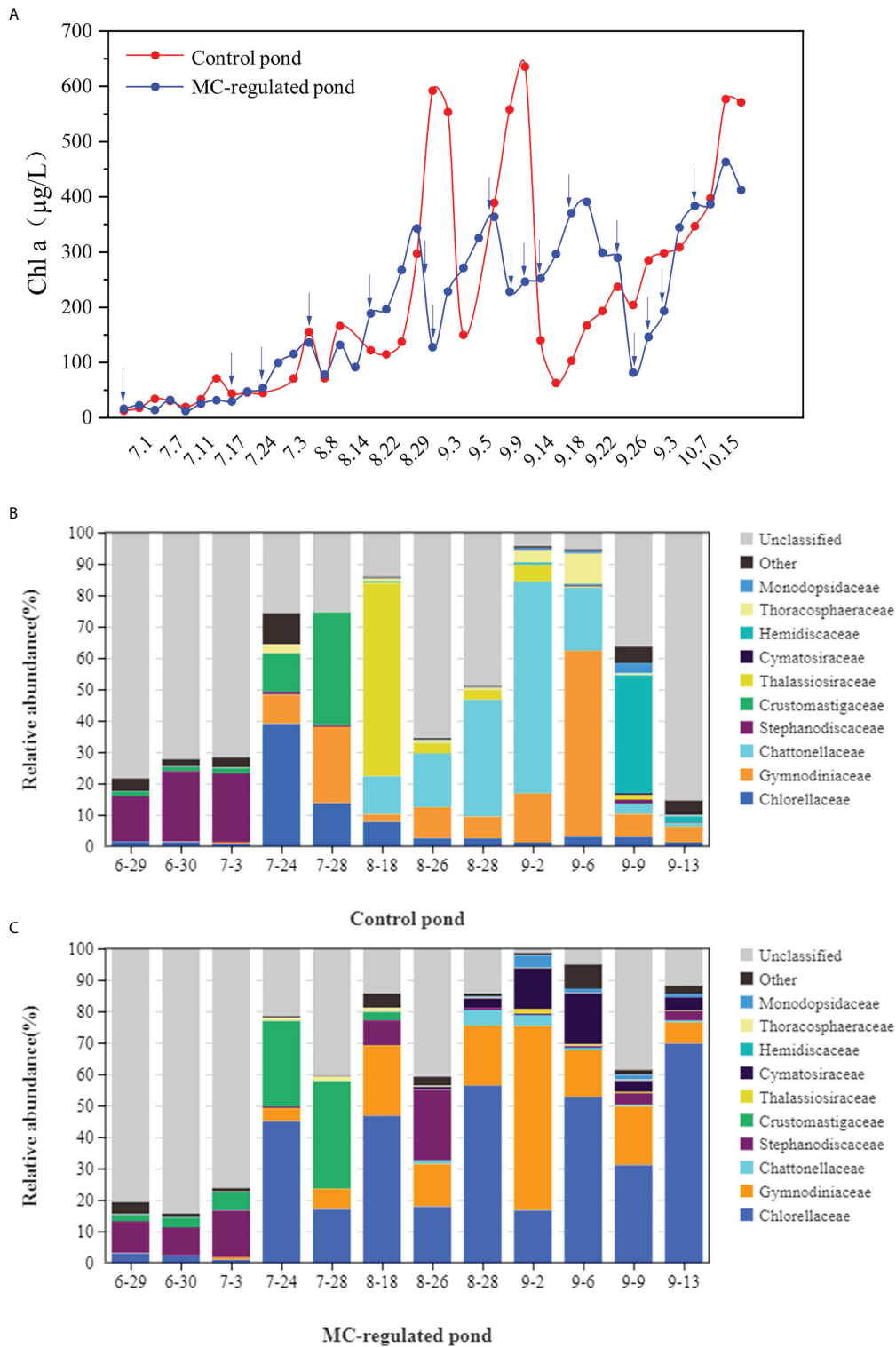


a decreasing ratio of diatoms was observed. Moreover, the dominant phytoplankton species exhibited different variation pattern between the two ponds. Notably, the dominant phytoplankton species in the control pond changed to dinoflagellates, mainly including Gymnodiniaceae at the family level. And the proportion of dinoflagellates (30.02%) in the control pond was higher than that of the MC-regulated pond (25.37%) (Welch's *t*-test,  $P > 0.05$ ). In addition, the Chattonellaceae (*Heterosigma* at the genus level), one of the HABs species, was another dominant family in the control pond, and had a much higher proportion in the control pond (13.09%) than the MC-regulated pond (0.92%). In contrast, the dominant phytoplankton species in the MC-regulated pond became green algae, mainly including Chlorellaceae at the family level and *Nannochloris* at the genus level, which had the highest proportion. In addition, the mean relative abundance (29.9%) of Chlorellaceae was significantly higher than that in the control pond (6.41%) (Welch's *t*-test,  $P < 0.05$ ).

### RDA ranking analysis between the abundance of major phytoplankton species and environmental variables

Overall, the phytoplankton richness, indicated by Chl *a* concentration, was negatively correlated with salinity and

temperature, and positively correlated with nutrients in both organic and inorganic forms (Figure 5), resulting in gradually higher phytoplankton biomass as the nutrient accumulated during the study period. As for the phytoplankton communities, the RDA results indicated that the phytoplankton patterns were largely grouped by nutrient regimes. In addition, there were temporal variations in the relationship between phytoplankton communities and environmental variables. In the initial stage, the clustering characteristics of the samples from the control and MC-regulated ponds were basically the same. During this period, the phytoplankton composition was dominated by diatoms, mainly including Stephanodiscaceae, whose abundance was positively correlated with DSi. Thereafter, the clustering characteristics of the samples from the two shrimp ponds differed, indicating that the environmental influences on the distribution of the samples from the two ponds became inconsistent. In the control pond, the dominant phytoplankton species have become harmful dinoflagellates, mainly including Gymnodiniaceae and Thoracosphaeraceae. And their abundance responded strongly to TON and TOP, as indicated by their positive correlations. In contrast, the dominant species of phytoplankton in the MC-regulated pond was Chlorellaceae, a kind of green algae, and its abundance exhibited a significant positive correlation with TOP and DIP and a significant negative correlation with DSi.



**FIGURE 4**  
 Temporal variations in the (A) Chl a concentrations and relative abundance of major taxa of phytoplankton in (B) the control pond and (C) MC-regulated pond at the family level. The blue arrows indicates the date when MC was applied.

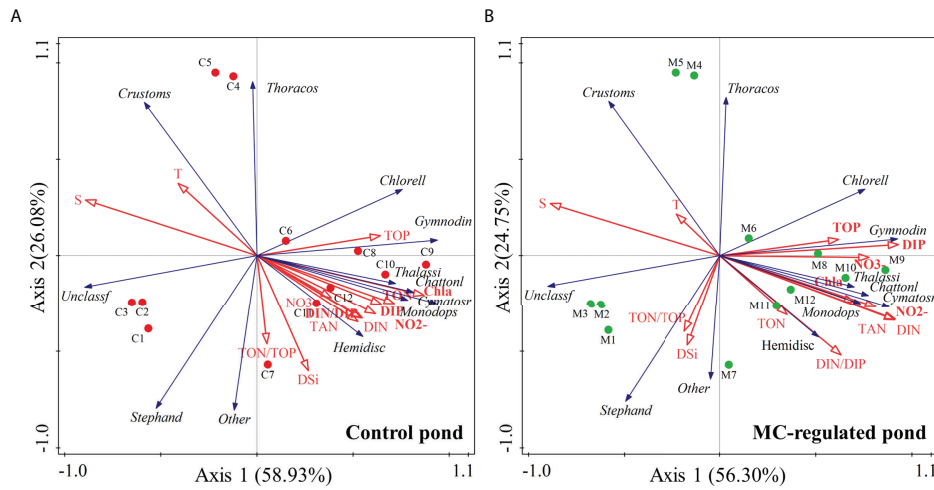


FIGURE 5 RDA ranking of samples and phytoplankton families with environmental variables in (A) the control pond and (B) the MC-regulated pond.

## Discussion

### Nutrient variation characteristics in mariculture ponds and their impacts on water quality of receiving coastal waters

In the intensive mariculture pond systems, the variation and accumulation of nutrients could cause the deterioration of water quality and pose threat to the organisms (Alonso-Rodriguez and Páez-Osuna, 2003; Casé et al., 2008; Ray et al., 2011; Yang et al., 2017; Díaz et al., 2019). In addition, the export of mariculture pond effluents is an important anthropogenic source of nutrients pollution in coastal waters (Cao et al., 2017; Li et al., 2017; Yang et al., 2017), which has significantly increased the nutrients concentrations (Paerl, 2006; Danielsson et al., 2008; Glibert and Burford, 2017), and altered the nutrients composition and structure (Glibert et al., 2012; Peñuelas et al., 2012; Sutton et al.,

2013). In the present study, the nutrients of the mariculture shrimp ponds exhibited unique characteristics compared with typical coastal regions. The concentrations of TON/P and DIN/P in the shrimp ponds were significantly higher, while DSi concentrations were lower than those in typical coastal waters. Notably, organic forms of nitrogen and phosphate were dominant (Figure 6), which could be attributed to continuous input and lower utilization efficiency of feeds (Bouwman et al., 2013). In general, more than 63% of nitrogen and 83% of phosphorus from feed have been discharged into the water column and sediment (Bouwman et al., 2013). Moreover, a large supply of organic nutrients stimulates microbial decomposition in the water column and sediment, as indicated by the increasing trend of DIN concentrations and DIN/TN ratios in this study. Notably, TAN was the predominant species of DIN, and it could be oxidized to  $\text{NO}_2^-$  and  $\text{NO}_3^-$ . Yang et al., 2017 found that the sediment fluxes in the shrimp ponds were

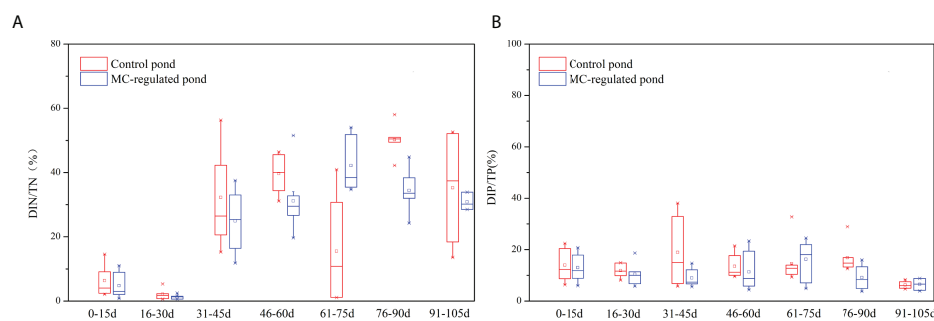


FIGURE 6 Temporal variations in (A) DIN/TN and (B) DIP/TP ratios.



significantly higher than the range of 0.01–31.25 mg N/(m<sup>2</sup> h). As the most common toxicant in mariculture (Santacruz-Reyes and Chien, 2010), the general safe concentrations for TAN and NO<sub>2</sub><sup>-</sup> in mariculture systems are under 200 µg/L and 10 µg/L, respectively (Lai, 2014), although the specific toxicity is species-dependent. In this study, the concentrations of TAN and NO<sub>2</sub><sup>-</sup> were higher than the safe concentrations, and this result was consistent with the study by Yang et al. (2017), which could potentially restrict the development of shrimp farming.

Moreover, as an important anthropogenic source of nutrient pollution in coastal waters (Table 1). (Lacerda et al., 2008) estimated that more than 827 t N/yr and 69.2 t P/yr were exported from shrimp ponds of northeastern Brazil. In the Gulf of California, the estimated nutrients loads from shrimp aquaculture were 9044 t N/yr and 3078 t P/yr (Páez-Osuna et al., 1997; Páez-Osuna et al., 2013; Páez-Osuna et al., 2017). In the present study, we further estimated the potential impacts of effluent discharge on water quality of receiving coastal waters. Assuming that our average data during the study period are representative of the aquaculture ponds across China, with a total area of 2.57×10<sup>10</sup> m<sup>2</sup> and a mean water depth of 1.4 m (Yang et al., 2017), approximately 1.05×10<sup>5</sup> t TON (in terms of N), 1.08×10<sup>4</sup> t TOP (in terms of P), 4.80×10<sup>4</sup> t DIN and 1.69×10<sup>3</sup> t DIP accumulated in the pond system and could be discharged into coastal waters. This value represents approximately 16% of the total nutrient fluxes from the main rivers of China into the sea, whose contribution was smaller than other anthropogenic nutrient sources (SOA (State Oceanic Administration), 2017). Our results were basically consistent with the estimation of Meng and Feagin, 2019, which proposed that more than 26% of the excess nitrogen in China's waters likely originates from shrimp production alone (Meng and Feagin, 2019). With the rapid development of marine aquaculture in China, its potential impact on water pollution will be more prominent in the future. In addition, the nutrients characteristics of the shrimp ponds could also significantly alter the composition and structure of

phytoplankton community and cause the occurrence of HABs, which could also limit healthy mariculture development.

## The impacts of nutrient variations on the phytoplankton community and potential occurrences of HABs

A healthy and stable phytoplankton community structure is crucial for the stability and balance of mariculture ecosystems (Alonso-Rodriguez and Páez-Osuna, 2003; Casé et al., 2008). Previous studies have documented the negative effect of algal blooms, especially dinoflagellate blooms, on shrimp development (Shumway et al., 1990; Alonso-Rodriguez and Páez-Osuna, 2003; Matsuyama and Shumway, 2009; Lou and Hu, 2014). Generally, the structure and variation of phytoplankton communities are controlled by the complex interactions between environmental drivers and biotic interactions (Griffiths et al., 2016). Among them, the concentration, composition, and structure of nutrients are crucial for the growth and community succession of phytoplankton and the potential occurrence of HABs. Generally, both the inorganic (Altman and Paerl, 2012; Kamp et al., 2015) and organic nutrients are available for phytoplankton (McCarthy, 1972; Lønborg and Álvarez-Salgado, 2012). This could also be verified by the positive correlations between the Chl *a* concentration with both the TON/P and DIN/P concentrations in the present study. Generally, most phytoplankton preferentially assimilate NH<sub>4</sub><sup>+</sup> due to the lower energy consumption requirements than those (Kamp et al., 2015). In addition, different species of phytoplankton differ in their preferences and responses to different forms of nutrients, previous studies indicated that NO<sub>3</sub><sup>-</sup> is preferred by diatoms (Goldman and Glibert, 1983; Lomas and Glibert, 1999; Lomas et al., 2002; Berg et al., 2003), while organic nitrogen is preferred by dinoflagellates (Glibert

TABLE 1 Main negative impacts caused by intensive mariculture systems in the Laizhou bay compared with other coastal areas.

Location of the mariculture systems	Main cultivated organism species	Main negative impacts	Reference
Laizhou bay, China	Shrimp	Increase in DIN/P, TON/P and Chl <i>a</i> concentrations, dinoflagellates bloom	This study
Weihai coastal area, China	Kelp, shellfish, and fish	Increase in DIN and DON concentrations	Li et al., 2017
Jiangsu coastal area, China	Seaweed, crab, and shellfish	Increase in DIN and DON concentrations, harmful macroalgal bloom	Liu et al., 2013
Hainan coastal area, China	Shrimp and fish	Increase in DIN and DON, dissolved organic carbon, and Chl <i>a</i> concentrations	Herbeck et al., 2013
Urias coastal lagoon, USA	Shrimp	Increase in nitrite and decrease in dissolved oxygen concentrations	Cardoso-Mohedano et al., 2016
Güllük Bay (Turkey)	Fish	Increase in inorganic N and P concentrations	Demirak et al., 2006
Coastal areas of the Gulf of California, USA	Shrimp	Contribution of 10.1% N and 3.3% P to total nutrient loading	Miranda et al., 2009

and Terlizzi, 1999; Dyhrman and Anderson, 2003; Fan et al., 2003). For instance, dinoflagellates could absorb DON and easily replace other algae as the dominant species when DON is the nitrogen source (Collos et al., 2014). In the present study, DON/P were the predominant species, although an increasing trend in the ratio of DIN was observed in the late stages, with an average proportion of 78.92% and 88.18% in TN and TP, respectively. Furthermore, the predominant TAN in DIN were observed in this study, with a significantly higher proportion than those of the  $\text{NO}_2^-$  and  $\text{NO}_3^-$ . This was consistent with previous studies, which could be attributed to the continuous decomposition of protein-rich feeds (Yang et al., 2017). Consequently, as the main forms of nutrient reservoirs, the nutrient pattern, dominated by TON and TAN, could contribute to the potential dominance of dinoflagellates, as verified by the 18S rDNA results.

In addition, the fluctuation in nutrient structure can induce a shift in dominant species of phytoplankton, and researchers have proposed that the variation of N/P ratios might stimulate HABs worldwide (Alonso-Rodriguez and Páez-Osuna, 2003; Davidson et al., 2012; Li et al., 2014). Generally, the pattern of “more N, less P and Si” may lead to a shift in the dominant species from diatoms to dinoflagellates (Li et al., 2014; Wang et al., 2015). For instance, excessive DIN and persistently elevated N/P have led to the dominant species shifting from diatoms to dinoflagellates in the Changjiang estuary (Wang and Cao, 2012). Likewise, Justić et al., (1995) suggested that the increase in N and P and the relative stabilization of Si in coastal waters, increased the possibility of Si limitation, leading to a shift in dominant phytoplankton species from diatom to non-diatom species. By comparing the differences in phosphorus requirements and uptake and utilization strategies of different algal species, the previous study has indicated that diatoms have the lowest mean optimum nitrogen to phosphorus ratio, followed by dinoflagellates, and green algae have the highest mean optimum nitrogen to phosphorus ratio (Hillebrand et al., 2013). In this study, there were no absolute concentration limitations of N, P, and Si in mariculture pond systems. However, the nutrient structure in terms of N/P and N/Si ratios exhibited significant temporal variations during the culture process. Relatively stable ratios of TON/TOP (with an average value of 27) but fluctuating increasing ratios of DIN/DIP (with an average value of 66) were found, which could be attributed to the higher mineralization and accumulation rate of N than P. In addition, increased DIN/DSi and decreased DSi/DIP ratios, indicating potential Si limitation, were observed especially in the middle and late stages. The variations in the nutrient stoichiometric ratios favored the shift in the dominant species that changed from diatoms to dinoflagellates. In addition, a gradually increasing proportion of dinoflagellates and decreasing proportion of diatoms corresponded well with the variations in N/P and N/Si ratios.

## Regulation effects of MC on nutrients and the phytoplankton community

The intensive mariculture systems are under the threat of excessive organic loading and nutrient accumulation, which cause water quality problems and subsequent diseases (Hargreaves and Tucker, 2004; Santacruz-Reyes and Chien, 2010; Castillo-Soriano et al., 2013; Hu et al., 2014). Notably, the TAN threat can become pronounced in intensive culture systems when TAN is rapidly accumulated to concentrations beyond the safe level (Santacruz-Reyes and Chien, 2010). More importantly, it can trigger outbreaks of HABs and pose a serious threat to the aquaculture ecosystem (Huang et al., 2016; Brown et al., 2020). In this study, the initiative MC technology was adopted to regulate the nutrients and phytoplankton in the typical mariculture pond system. We observed that the nutrient contents of both organic and inorganic forms in the water column effectively decreased in most instances 24 h after the spray of MC (Figure 7). The TON and TOP contents could be reduced by up to 57% and 65%, respectively. While the TAN,  $\text{NO}_2^-$  and  $\text{PO}_4^{3-}$  concentrations could be reduced by up to 32%, 45%, and 64%, respectively, in the MC-regulated pond. In contrast, both organic and inorganic forms in the water column of the control pond increased in most instances. Overall, the contents of these nutrients in the MC-regulated pond were lower than those in the control pond. It has been suggested that MC can reduce inorganic nutrients, especially phosphate, and organic nutrients through adsorption flocculation and chelation (Lu et al., 2015; Yu et al., 2017). In addition, MC could cause algal cells to flocculate and settle to the bottom layer, and the interaction of clay minerals with organic matter provides physical protection for organic matter (Hemingway et al., 2019). This protection could delay the microbial (Pinck and Allison, 1951) and oxidative (Eusterhues et al., 2003) decomposition processes of organic matter and reduce the mineralized regeneration of nutrients.

Furthermore, MC exerted a moderating effect on the phytoplankton biomass and community in the shrimp pond. In this study, the phytoplankton biomass was significantly higher and exhibited greater volatility in the control pond. In addition, HABs have occurred twice in the control pond, including an *H. akashiwo* bloom on September 2 with a density of  $1.26 \times 10^5$  cells/mL (He et al., unpublished data). In contrast, the phytoplankton community in the MC-regulated pond was relatively stable in change and no HABs occurred during the study period. Based on the mean Bray-Curtis distances and molecular ecological network analysis of the community, (Ding et al. 2021) proposed that MC could enhance the resistance of phytoplankton ecological communities in cultured waters to environmental disturbance. Moreover, under the traditional condition, a high propensity for dinoflagellate blooms existed, influenced by the variation in the

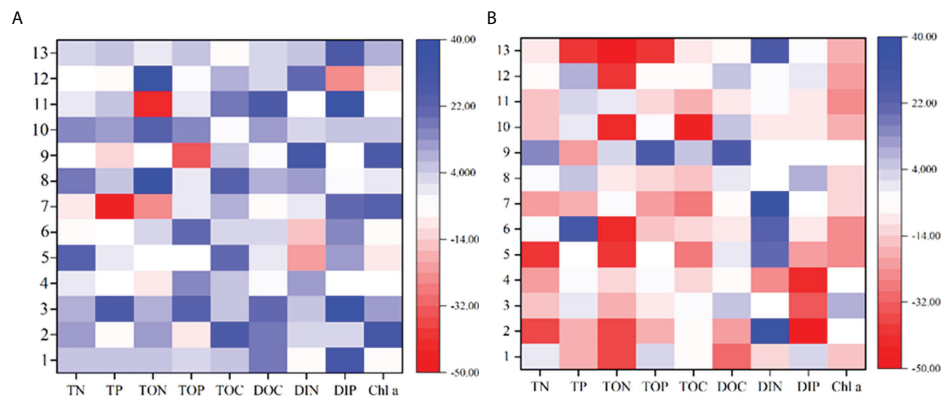


FIGURE 7  
Changes in nutrients and Chl a in (A) the MC-regulated pond before and after spraying of MC and the comparison with (B) the control pond.

composition and structure of nutrients, and posed a potential risk to the growth of shrimp. This result could be verified by the variation in the dominant species of phytoplankton that changed from diatoms to dinoflagellates in the control pond. Compared with the control pond, the percentage of dinoflagellates in the MC-regulated pond was maintained at a lower level, and the percentage of green algae was higher, with *Nannochloris* being the dominant species. As an aquatic bait algae species, *Nannochloris* can improve the survival and production rate of fish and shrimp to some extent, which is important to maintain the stability of the ecosystem (Alonso-Rodriguez and Páez-Osuna, 2003; Davidson et al., 2009). MC can directly dispose of HAB organisms through flocculation; additionally, they can induce the programmed mortality of red tide organisms through oxidative stress and other effects, thus controlling HABs (Yu et al., 2017). On the one hand, the *Nannochloris* could occupy a favorable ecological niche and became dominant after the removal of targeted dinoflagellates in the MC-regulated pond. On the other hand, the MC increased the N/P ratio and favored the *Nannochloris*, which has a higher average optimal nitrogen to phosphorus ratio than dinoflagellates (Hillebrand et al., 2013).

## Conclusion

This research studied the dynamic variations in nutrients and phytoplankton of intensive mariculture systems and explored the effects of MC. The intensive culture of *Litopenaeus vannamei* caused a temporal significant increase in nutrients, especially in the organic forms. In addition, concurrently with ascending N/P ratio and decreasing Si/N and Si/P ratios, a marked increase in the biomass and ratios of dinoflagellates to diatoms abundance were also observed, which pose a potential threat to the mariculture organism. The MC reduced the contents of nutrients in both organic and inorganic forms, and improved the water quality.

Moreover, MC effectively removed the dinoflagellates and contribute to the dominance of *Nannochloris*, which improved the stability of the phytoplankton community. This study provide new insights into an effective regulation treatment for managing water quality and maintaining sustainable mariculture development.

## Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

## Author contributions

LC: writing – original draft and investigation. YD: writing – review & editing and formal analysis. LH: writing – review & editing and data curation. ZW: writing – review & editing and data curation. YY: writing – review & editing and investigation. XC: writing – review & editing and formal analysis. XS: writing – review & editing. ZY: writing – review & editing and supervision. All authors contributed to the article and approved the submitted version.

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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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## Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2022.976353/full#supplementary-material>

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