

## Cyanobacteria reduction in the integrated culture of Pacific white shrimp and West India pointed venus in a biofloc system

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### Abstract

This study aimed to evaluate the integrated culture of the Pacific white shrimp (*Litopenaeus vannamei*) and the West Indian pointed venus (*Anomalocardia flexuosa*) reared in a biofloc system. The performance of both species and the phytoplankton community were evaluated. Shrimp monoculture and three stocking densities of West Indian pointed venus (2.5, 5.0 and 7.5 Kg m<sup>-3</sup>) were evaluated in 0.05 m<sup>3</sup> tanks. First of all, the addition of *A. brasiliiana* did not affect the water quality parameters of the culture. Regarding the phytoplankton community, 13 taxa were found: 5 cyanobacteria, 4 dinoflagellates, 3 diatoms and 1 green algae. Although diatoms were dominant at the beginning of cultivation, in the middle and at the end a cyanobacterial dominance was observed (proven by ANOSIM and nMDS). The taxa that most contributed to the abundance of cyanobacteria were *Aphanocapsa* sp. and *Pseudanabaena* sp. Regarding the performance of the shrimp, all integrated treatments present the higher growth performances (final weight, yield and survival) when compared to shrimp monoculture. Moreover, gradual mortalities of *A. brasiliiana* were observed, probably due to the high concentration of cyanobacteria. Thus, it can be observed that the integration of molluscs in shrimp culture can reduce the concentration of harmful algae and improve the zootechnical performance of the shrimp.

Keywords: *Anomalocardia flexuosa*, harmful algae, Integrated Multitrophic Aquaculture, *Litopenaeus vannamei*, water quality.

### INTRODUCTION

Biofloc systems have been used to reduce the water exchange and consequently to reduce the wastewater production, in addition, it also recycles part of the metabolite waste from fish and shrimp nutrition. This makes the biofloc systems environmentally friendly because they reduce impacts and manage to have greater productivity compared to traditional systems (Crab *et al.*, 2012; Krummenauer *et al.*, 2012).

Biofloc uses an organic carbon source to maintain a satisfactory carbon:nitrogen (C:N) ratio that favors the growth of heterotrophic, chemoautotrophic and/or nitrifying bacteria. This bacteria community will assimilate the nitrogen compounds in the system, converting it into microbial biomass

(Ebeling *et al.*, 2006; Avnimelech, 2009). Molasses and cane sugar (sucrose) are the most widely used organic carbon sources due to their high availability and low cost, compared to the of organic carbon sources used for bacterial cultures and other heterotrophic and/or mixotrophic microorganisms (Teclu *et al.*, 2009, Lorenzo *et al.*, 2015).

In biofloc systems, there is a naturally organic matter accumulation (Krummenauer *et al.*, 2011), which can favor the growth of other microorganisms harmful to species in cultivation as: *Vibrio* spp. (Ferreira *et al.*, 2011; Manan *et al.*, 2017) and harmful algae (Brito *et al.*, 2014). To remedy this problem, integrated multi-trophic aquaculture (IMTA) systems have been used to reduce organic waste and also the nitrogen and phosphorus compounds levels. IMTAs have

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been an efficient strategy to minimize the environmental problems associated with the monoculture system (Angel & Freeman, 2009).

Among the organisms with potential for IMTA are: algae, fish, crustaceans and bivalve molluscs (Troell *et al.*, 2009). Bivalve molluscs are attractive because they reduce the organic matter of the farming systems while the fish can act in phosphorus recovery through its use for skeletal formation (which is absent in crustaceans) (Oliveira & Brito, 2005; Modesto *et al.*, 2010; Jones *et al.*, 2002).

The residues accumulation in biofloc may influence the phytoplankton community, due to eutrophication (Boyd, 2015), which may result in an increase in microalgal biomass, and a reduction in the diversity of planktonic organisms, with a predominance of harmful algae (Hu *et al.*, 2018), harming the production of the target organism of the culture (Boyd, 2015; Gao *et al.*, 2017).

Many species of harmful algae are potential toxins producers, and for this reason, their abundance raises concerns due to the reduction of oxygen concentrations, in the dark period, and risks to cultivated organisms as well as to their consumers. Cyanobacteria and dinoflagellates are the most harmful algae groups in aquaculture systems; on other hand, diatoms and green algae are the most desired algae groups in aquaculture systems (Sant'Anna & Azevedo, 2000; Baudin *et al.*, 2006; Van Apeldoorn *et al.*, 2007). The presence of toxic cyanobacteria has already been reported in intensive (Brito *et al.*, 2014; Marinho *et al.*, 2014) and semi-intensive (Barbosa, 2015) shrimp farming.

In this way, the first goal of this study was to evaluate the co-culture of two species of important economic interest, Pacific white shrimp (*Litopenaeus vannamei*) and West Indian pointed venus (*Anomalocardia flexuosa*) integrated in a biofloc system. In addition, this study sought to evaluate the ecological influence of *A. flexuosa* at three different densities in the phytoplankton community.

## MATERIALS AND METHODS

### *Biological material*

*L. vannamei* post-larvae ( $2.73 \pm 0.25$  mg) were purchased from a commercial laboratory (Aquatec LTDA). *A. flexuosa* specimens were collected at Mangue Seco beach, Igarassu, Pernambuco, Brazil (07° 49' 44.19''S, 035° 50' 03.06''W). This work does not require approval from the ethics committee on animal use.

### *Experimental design and conditions*

The experiment was conducted for 42 days at the Laboratório de Maricultura Sustentável (LAMARSU), which is part of the Departamento de Pesca e Aquicultura at the Universidade Federal Rural de Pernambuco (UFRPE). The experiment was performed in a completely randomized design with four replicates, integrating shrimp (*L. vannamei*) with a bivalve mollusc (*A. flexuosa*) reared in a biofloc system.

Five days before the storage of shrimp and molluscs, water from a biofloc matrix tank (total ammonia 0.25 mg L<sup>-1</sup>, nitrite 1.75 mg L<sup>-1</sup>, alkalinity 107.00 mg CaCO<sub>3</sub> L<sup>-1</sup>, pH 8.30, 30 salinity and sedimentable solids 8 mL L<sup>-1</sup>), was homogenized and uniformly distributed to fill two tanks.

The experimental units consisted of a 0.05 m<sup>3</sup> tank. No water exchange occurred during this time, freshwater was only used to replace water loss due to evaporation. Ammonia control was performed daily by adding molasses as an organic carbon source according to Avnimelech (1999). Moreover, hydrated lime (Ca(OH)<sub>2</sub>) was added in 10% (by weight) of the daily feed supply, throughout the experimental period, in order to maintain the alkalinity and pH above 100 mg L<sup>-1</sup> and 7.5, respectively. Three cylindrical porous stones at each experimental unit were used to keep the bioflocs in suspension and maintain oxygen above 5 mg L<sup>-1</sup>. The luminous intensity was maintained at 2000 lux with a 12h photoperiod.

Shrimps were fed with a 40% crude protein commercial feed (Guabi-Potimirim) four times a day according the Van-Wyk (1999) feed table. In addition, they were reared at a density of 3,000 shrimp m<sup>-3</sup>. The experiment consisted of four treatments corresponding to a shrimp monoculture (BFT) and three stocking densities of *A. flexuosa*: 2.5 (BFT 2.5), 5.0 (BFT 5.0) and 7.5 (BFT 7.5) Kg wet weight m<sup>-3</sup>. The mollusc specimens were washed and sanitized with a sodium hypochlorite bath at 5 g L<sup>-1</sup> for five minutes, then they were acclimated to the experimental conditions in 0.35 m<sup>3</sup> tanks with seawater (30 salinity), at 26 °C and a density of 3 molluscs L<sup>-1</sup>. The acclimatization of the molluscs occurred for five days before they were inserted in the experimental units, where 20 L of water from the biofloc system were added daily.

### *Water quality*

Dissolved oxygen, salinity, temperature, total dissolved solids and pH were measured twice a day with a multiparameter probe (YSI model 556, Yellow Springs, Ohio, USA). Total ammonia (mg L<sup>-1</sup>), nitrite (mg L<sup>-1</sup>) and alkalinity (mg CaCO<sub>3</sub> L<sup>-1</sup>) were measured every two days using the colorimetric test (Labcon Test).

### *Phytoplankton community*

For evaluation of the phytoplankton community, water samples were collected at the beginning, middle (21st day) and end (42th day) of the experiment. To do so, 500 mL samples of water from each experimental unit were filtered through a cylindrical net with 90 µm mesh in order to decrease the suspended solids amount in the sample, then they were filtered with a 50 µm mesh to remove the zooplankton, and finally 15 µm for phytoplankton collection. The filtered material was immediately fixed in 4% formalin and stored for further analysis. The identification and quantification of phytoplankton samples were performed using a Sedgewick-Rafter chamber and binocular optical microscope (Olympus, CH30) with a 1.000 x magnification. The organisms were identified based on

cell morphology at the lowest possible taxonomic level with the aid of specific identification keys (Bicudo & Menezes, 2006; Wehr *et al.*, 2015). To calculate the abundance, the methodology described by Hötzel & Croome (1999) was followed. Besides, the groups were categorized according to the relative abundance classification, which they consider in rare (<10%), less abundant (10%-40%), abundant (40%-70%) and dominant (>70%) (Neumann-Leitão *et al.*, 1995).

### Shrimp and mollusc performance

Shrimp performance was evaluated by weekly biometry. At the end of rearing, survival (%), final mean weight (g), specific growth rate (SGR), biomass gain (g) and yield (Kg m<sup>-3</sup>) were measured. Mollusc survival was also evaluated weekly.

### Statistical analysis

All data are presented with mean  $\pm$  standard deviation (n = 3). First, the homogeneity of the variances (Cochran) and normality (Shapiro-Wilk) were evaluated. Subsequently, an ANOVA was applied, followed by the Tukey's means comparison test, when necessary, using a level of 0.05 (Zar, 1999). Moreover, cluster analysis (Bray-Curtis similarity) and a non-metric multidimensional scaling (nMDS) were carried out, aiming to observe the similarity of the community in temporal and spatial scale, and their possible group formation; finally, an ANOSIM, with 999 permutations, in order to identify the difference between groups of phytoplankton (Clarke, 1993). PRIMER 6.0 and Assistat 7.6 software's were used.

## RESULTS

### Water quality

Data on water quality variables are shown in Table 1. There were no significant differences ( $p \geq 0.05$ ) between water quality parameters, except for alkalinity ( $p < 0.05$ ). During the experiment, the dissolved oxygen and temperature

remained close to 5 mg L<sup>-1</sup> and 27 °C, respectively. The pH and alkalinity remained stable near to 8.0 and 120 mg CaCO<sub>3</sub> L<sup>-1</sup>, respectively.

### Phytoplankton community

A total of 13 taxa were identified: 5 cyanobacteria, 3 diatoms, 4 dinoflagellates and 1 green algae, and are listed in Table 2. The richness estimate was 13 for the Chao and Jackknife samplers, thus, the richness estimators reached the asymptote with the observed taxa number. At the beginning of the cultivation, the most representative algae group was diatoms (73.68%), represented mainly by *Thalassiosira* spp. In the middle and final of culture, cyanobacteria (ranging from 97.98 to 99.98%) were dominant in all treatments, reaching a maximum density of 69,640.5 cells. mL<sup>-1</sup> at the 42<sup>th</sup> day of culture in the BFT treatment (shrimp monoculture). During that same period diatoms were considered rare, while the dinoflagellates and green algae were rare throughout the experimental period (Figure 1).

In the treatments of integrated culture, the lowest density of cyanobacteria was observed, differing significantly ( $p < 0.05$ ) from the BFT treatment, both in the middle and at the end of the experiment. In the middle of culture (21 days), the shrimp monoculture showed from 4.37 to 7.00 times more cyanobacteria than the treatments with *A. flexuosa*. At the end of culture, this value was 47.69, 17.77 and 113.14 times higher, for BFT 2.5, BFT 5 and BFT 7.5, respectively.

The ANOSIM did not show significant differences in the phytoplankton community among the treatments (Global R = -0.277), but it showed that in relation to the days of cultivation, phytoplankton showed a significant difference (Global R = 0.944).

There is a 50% cut, the cluster showed the formation of 2 groups: the group I was composed by day 0 (beginning of the experiment), and the group II was composed by 21 and 42 days - showed a similarity of 61.36% (Figure 2A). The nMDS analysis confirmed this differentiation of the phytoplankton

Table 1. Water quality parameters in the integrated culture of Pacific white shrimp and West India pointed venus in a biofloc system.

Parameter	Treatments			
	BFT	BFT 2.5	BFT 5.0	BFT 7.5
Temperature (°C)	26.47 $\pm$ 0.07 <sup>a</sup>	26.40 $\pm$ 0.07 <sup>a</sup>	26.40 $\pm$ 0.08 <sup>a</sup>	26.45 $\pm$ 0.07 <sup>a</sup>
Dissolved oxygen (mg L <sup>-1</sup> )	4.72 $\pm$ 0.05 <sup>a</sup>	4.78 $\pm$ 0.49 <sup>a</sup>	4.76 $\pm$ 0.06 <sup>a</sup>	4.73 $\pm$ 0.05 <sup>a</sup>
Salinity	33.09 $\pm$ 0.46 <sup>a</sup>	33.20 $\pm$ 0.38 <sup>a</sup>	34.00 $\pm$ 0.53 <sup>a</sup>	33.51 $\pm$ 0.46 <sup>a</sup>
TDS (g L <sup>-1</sup> )	32.80 $\pm$ 0.41 <sup>a</sup>	33.04 $\pm$ 0.37 <sup>a</sup>	33.61 $\pm$ 0.46 <sup>a</sup>	33.25 $\pm$ 0.42 <sup>a</sup>
pH	8.14 $\pm$ 0.04 <sup>a</sup>	8.18 $\pm$ 0.03 <sup>a</sup>	8.15 $\pm$ 0.04 <sup>a</sup>	8.17 $\pm$ 0.03 <sup>a</sup>
Ammonia (mg L <sup>-1</sup> )	0.25 $\pm$ 0.03 <sup>a</sup>	0.44 $\pm$ 0.08 <sup>a</sup>	0.79 $\pm$ 0.34 <sup>a</sup>	0.40 $\pm$ 0.14 <sup>a</sup>
Nitrite (mg L <sup>-1</sup> )	1.96 $\pm$ 0.31 <sup>a</sup>	2.00 $\pm$ 0.36 <sup>a</sup>	1.73 $\pm$ 0.35 <sup>a</sup>	2.62 $\pm$ 0.17 <sup>a</sup>
Alkalinity (mg L <sup>-1</sup> )	133.30 $\pm$ 9.00 <sup>ab</sup>	115.40 $\pm$ 8.50 <sup>b</sup>	118.60 $\pm$ 10.11 <sup>ab</sup>	146.20 $\pm$ 9.71 <sup>a</sup>

Data presented as mean  $\pm$  standard deviation (n = 3). Different letters on the same line indicate a significant difference ( $p < 0.05$ ) using the Tukey test. TDS- Total dissolved solids.

Table 2. Density (cells mL<sup>-1</sup>) and average relative abundance (%) of phytoplankton taxa in the integrated culture of Pacific white shrimp and West India pointed venus.

Taxa		Beginning	Middle (21st day)				End (42th day)			
		BFT	BFT 2.5	BFT 5.0	BFT 7.5	BFT	BFT 2.5	BFT 5.0	BFT 7.5	
Diatoms	(cells mL <sup>-1</sup> )	164.3	8 <sup>a</sup>	14.2 <sup>a</sup>	22.6 <sup>a</sup>	2.3 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>
	(%)	73.67	0.01	0.07	0.11	0.02	0	0	0	0
<i>Thalassiosira</i> sp.1	(cells mL <sup>-1</sup> )	107	7	11.6	18.6	2.3	0	0	0	0
	(%)	46.38	0.01	0.06	0.09	0.02	0	0	0	0
<i>Thalassiosira</i> sp.2	(cells mL <sup>-1</sup> )	56	1	2.6	4	0	0	0	0	0
	(%)	26.79	0	0.01	0.02	0	0	0	0	0
<i>Pseudostaurosira</i> sp.	(cells mL <sup>-1</sup> )	1.3	0	0	0	0	0	0	0	0
	(%)	0.5	0	0	0	0	0	0	0	0
Cyanobacteria	(cells mL <sup>-1</sup> )	48.6	56,239.5 <sup>a</sup>	12,808.7 <sup>b</sup>	12,013.6 <sup>b</sup>	8,024.9 <sup>b</sup>	69,491.5 <sup>a</sup>	1,457.2 <sup>b</sup>	3,910.4 <sup>b</sup>	614.2 <sup>c</sup>
	(%)	20.08	99.91	99.84	99.81	99.82	99.98	99.65	99.9	97.98
<i>Pseudanabaena</i> sp.	(cells mL <sup>-1</sup> )	21.3	10.3	951.3	12	11.6	52,016.6	287	1,217.3	188.9
	(%)	7.52	0.02	13.38	0.15	0.15	65.48	36.66	46.84	57.49
<i>Synechococcus</i> sp.1	(cells mL <sup>-1</sup> )	24.3	0	0	0	0	0	0	0	0
	(%)	11.75	0	0	0	0	0	0	0	0
<i>Synechocystis</i> sp.2	(cells mL <sup>-1</sup> )	3	0	0	0	0	0	0	0	0
	(%)	0.81	0	0	0	0	0	0	0	0
<i>Chroococcus</i> sp.	(cells mL <sup>-1</sup> )	0	417	182.6	159	67	218.3	124.3	958.6	46
	(%)	0	6.06	1.86	1.83	0.87	1.47	10.54	16.25	13.12
<i>Aphanocapsa</i> sp.	(cells mL <sup>-1</sup> )	0	52,052.20	11,674.80	11,843	7,946.3	17,256.60	1,045.9	1,734.50	379.3
	(%)	0	93.83	84.6	97.83	98.8	33.03	52.45	36.81	27.37
Dinoflagellates	(cells mL <sup>-1</sup> )	11.2	33.9 <sup>a</sup>	17.2 <sup>a</sup>	7.6 <sup>a</sup>	13.9 <sup>a</sup>	6.3 <sup>a</sup>	3.6 <sup>a</sup>	3.3 <sup>a</sup>	2.3 <sup>a</sup>
	(%)	6.05	0.07	0.1	0.07	0.15	0.01	0.35	0.09	2.02
<i>Schuetiella mitra</i>	(cells mL <sup>-1</sup> )	1.3	0.3	2.3	0.3	1	0	0	0	0
	(%)	0.66	0	0.01	0	0.01	0	0	0	0
<i>Oxyrrhis</i> sp.	(cells mL <sup>-1</sup> )	5	4.6	2.6	1.3	2.6	0.3	0	0	0
	(%)	1.83	0.01	0.02	0.02	0.03	0	0	0	0
Lophodinium sp.	(cells mL <sup>-1</sup> )	0.6	0	0	0	0	0	0	0	0
	(%)	0.16	0	0	0	0	0	0	0	0
<i>Coolia</i> sp.	(cells mL <sup>-1</sup> )	4.3	29	12.3	6	10.3	6	3.6	3.3	2.3
	(%)	3.4	0.06	0.07	0.05	0.11	0.01	0.35	0.09	2.02
Green algae	(cells mL <sup>-1</sup> )	0.3 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>	0 <sup>a</sup>
	(%)	0.18	0	0	0	0	0	0	0	0
<i>Diplochloris</i> sp.	(cells mL <sup>-1</sup> )	0.3	0	0	0	0	0	0	0	0
	(%)	0.18	0	0	0	0	0	0	0	0
Total	(cells mL <sup>-1</sup> )	224.4	56,281.4	12,840.1	12,043.8	8,041.1	69,646.8	1,460.8	3,913.7	616.3

Data presented as mean ± standard deviation (n = 3). Different letters on the same line indicate a significant difference (p < 0.05) using the Tukey test.

community, over time, through the distance between the groups formed (Figure 2B).

The SIMPER multivariate analysis showed at beginning, middle and end of the culture, the typified

taxa were *Thalassiosira* sp.1 (36.89%), *Chroococcus* sp. (31.09%) and *Chroococcus* sp. (45.07%), respectively. Furthermore, with this analysis it was found that in relation to treatments throughout the experimental period, the

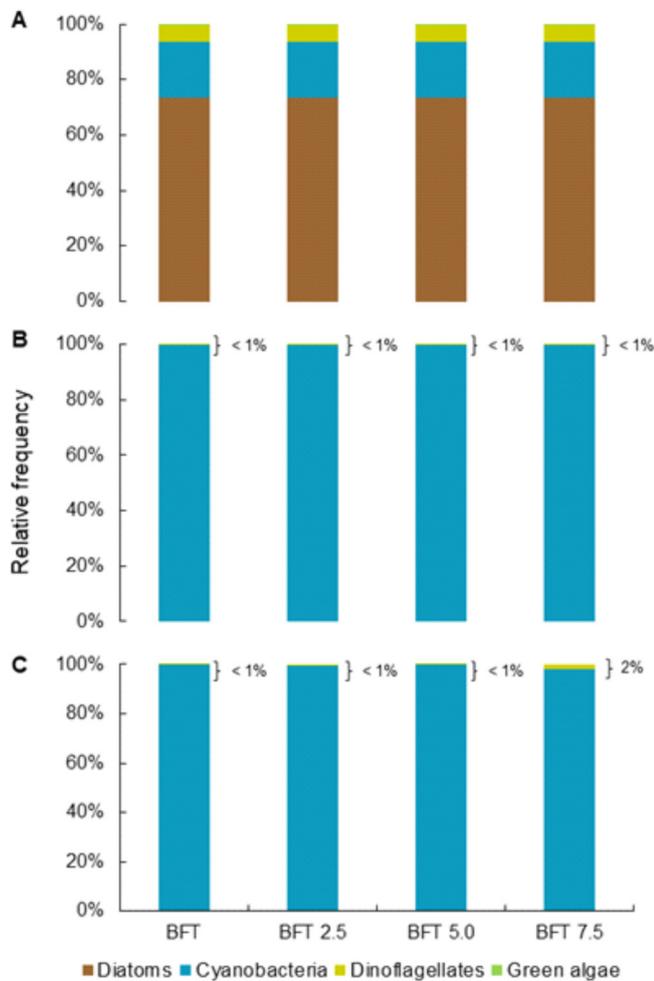


Figure 1. Relative abundance of phytoplankton presents in the Pacific white shrimp and West Indian pointed venus integrated in a biofloc system at 0 (A), 21 (B) and 42 (C) days of cultivation.

taxa *Pseudanabaena* sp. (36.03 – 65.83%) was the most contributed in all treatments.

**Zootechnical performance**

At the end of the 42 days of the culture, was observed that the shrimp performance was similar among the treatments of integrated cultivation *A. flexuosa*, presented a higher final weight, survival, biomass gain and productivity when compared to monoculture (Table 3). Nevertheless, a high mortality of *A. flexuosa* in all stocking densities used in the present study (Figure 3).

**DISCUSSION**

The water quality variables were kept within the ideal range for the cultivation of the Pacific white shrimp (Samocha *et al.*, 2017). The low nitrogen compounds suggested that ammonia control was mostly driven by heterotrophic pathway (Poli *et al.*, 2019). Temperature and salinity were also according to life environment of West Indian pointed venus (Oliveira *et al.*, 2011) and also culturing systems (Oliveira *et al.*, 2019a).

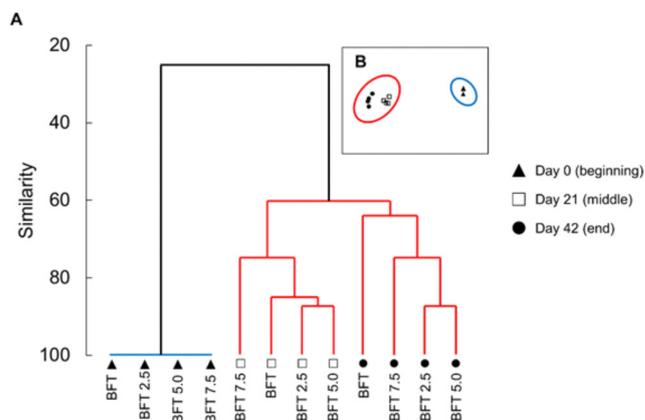


Figure 2. Cluster analysis (A) and non-metric multidimensional scaling (nMDS) (B) of the phytoplankton community present in the Pacific white shrimp and West Indian pointed venus integrated in a biofloc system.

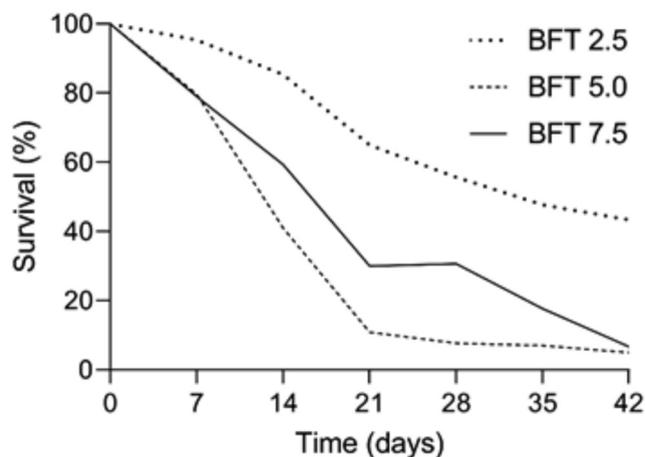


Figure 3. Survival (%) of West Indian pointed venus integrated with Pacific white shrimp in a biofloc system.

Table 3. Zootechnical performance of Pacific white shrimp in the integrated culture with West India pointed venus in a biofloc system.

Parameter	Treatments			
	BFT	BFT 2.5	BFT 5.0	BFT 7.5
Weight (g)	0.15±0.02 <sup>b</sup>	0.31±0.06 <sup>a</sup>	0.25±0.01 <sup>a</sup>	0.27±0.05 <sup>a</sup>
Survival (%)	38.56±3.28 <sup>b</sup>	58.89±3.31 <sup>a</sup>	61.33±1.15 <sup>a</sup>	62.56±4.13 <sup>a</sup>
Biomass gain (g)	8.63±1.31 <sup>b</sup>	25.08±3.6 <sup>a</sup>	21.31±1.21 <sup>ab</sup>	24.61±5.50 <sup>a</sup>
Productivity (Kg m <sup>-3</sup> )	0.17±0.03 <sup>b</sup>	0.50±0.07 <sup>a</sup>	0.43±0.02 <sup>ab</sup>	0.49±0.11 <sup>a</sup>
SGR	6.79±0.27 <sup>a</sup>	8.07±0.43 <sup>a</sup>	7.68±0.08 <sup>a</sup>	7.82±0.39 <sup>a</sup>

Data presented as mean ± standard deviation (n = 3). Different letters on the same line indicate a significant difference (p <0.05) using the Tukey test. SGR - specific growth rate.

From the third week of the cultivation on, a change in the culture watercolor was observed (typically brown color traditional biofloc systems, to a blue-green color), which motivated us to examine the phytoplankton community.

Diatoms were more abundant at the beginning of cultivation in all treatments, but over time there was a reduction in this abundance that may be linked to the gradual eutrophication of the system water, caused by the feeding rates and accumulation of organic matter that can favor groups of algae more resistant, such as cyanobacteria. Cyanobacteria have several biochemical, physiological, genetic and reproductive adaptations for the formation of blooms in adverse conditions (e.g., low light, production of info-chemicals, tolerance to high levels of nitrogenous and phosphorus compounds etc.) (Reichwaldt & Ghadouani, 2012; Yusoff *et al.*, 2010). Although dinoflagellates were not representative for this study, they are also known to have large impacts on aquaculture, such as the recent Godzilla Red Tide event in Chile, the largest fish farm mortality, equivalent to an export loss of USD\$ 800 million which when combined with shellfish toxicity, resulted in a major social-economic imbalance (Mardones, 2020; Trainer *et al.*, 2020). According to Yusoff *et al.* (2010), a diatom density reduction ( $9.19 \times 10^3$  to  $3.07 \times 10^3$  cells mL<sup>-1</sup>) was observed in a cyanobacterial bloom ( $7.20 \times 10^3$  to  $122.37 \times 10^3$  cells mL<sup>-1</sup>) during the shrimp farming in a eutrophic water.

In biofloc systems, there is a low transparency of the water, due to microbial flocs, and an increase in phosphorus compounds throughout culture mainly due to the decomposition of the uneaten feed and the excretion of the cultivated organisms. These facts are already exhaustively known for causing harmful cyanobacterial blooms (Oliveira *et al.*, 2019b) and they are the most likely causes for the dominance of cyanobacteria in the present study. Moreover, although there has been no negative impact on water quality parameters, it is also likely that low survival rates are due to toxin production (Kangur *et al.*, 2005; Malbrouck & Kestemont, 2006).

The most abundant cyanobacteria taxa found in this study, are potential toxin producers: *Aphanocapsa* genus is potential producers of microcystins. Cyanobacterial dominance already reported in a number of studies (Brito *et al.*, 2014; Wei *et al.*, 2016; Marinho *et al.*, 2017). According to Brito *et al.* (2014) and Marinho *et al.* (2017) a low nitrogen: phosphorus ratio can favor the cyanobacteria in aquaculture systems. This is because some cyanobacteria genera can to fix atmospheric nitrogen (N<sub>2</sub>) when in low concentrations in water. Whereas the bacterial community is well established in biofloc systems and it has a faster growth rate than algae and cyanobacteria (an order of magnitude), the atmospheric nitrogen assimilation is the best reason for the dominance of cyanobacteria in biofloc systems. Environmental factors such as light availability, temperature, dissolved inorganic compounds concentrations can result in a succession of dominant phytoplankton species (Casé *et al.*, 2008; Oliveira *et al.*, 2020) and this can explain the difference in dominance groups at the beginning, middle and end of culture.

The highest shrimp zootechnical performance was found in integrated cultivations (BFT 2.5, BFT 5.0 and BFT 7.5) when compared to shrimp monoculture. It is likely that the

increase in shrimp zootechnical performance was due to reducing the concentration of cyanobacteria cells by molluscs. This fact also justifies the low survival of molluscs that were probably infected by toxins (Queiroga *et al.*, 2017). Therefore, it is possible to affirm the following cascade effect: (i) West Indian pointed venus reduced the cyanobacterial load, which resulted in a (ii) highest zootechnical performance for shrimp in the integrated cultures. (iii) On the other hand, West Indian pointed venus specimens were probably intoxicated by cyanobacterial toxins. Molluscs, being filter organisms, can retain suspended particles in water and can function as a biological filter (Oliveira & Oliveira, 2019). This filtration can affect the amount of material present in the water column (Deslous-Paoli *et al.*, 1992), as happened with cyanobacteria.

Although the integrated shrimp and mollusc cultures have shown better zootechnical performance when compared to shrimp monoculture, other studies have already reported higher final weight (0.68 - 1.08 g), productivity (1.21 - 2.46 Kg m<sup>-3</sup>) and survival (71.3 - 95.6%) for Pacific white shrimp monoculture in biofloc systems. Moreover, a survival of West Indian pointed venus post-larvae of 53.24% was reported by Oliveira *et al.* (2019a) in a super-intensive culture under a recirculation aquaculture system. In addition to the higher concentration of cyanobacteria, shear forces and the high content of organic matter in biofloc systems may also have negatively influenced mollusc mortality (Rodrigues, 2009). Therefore, it is still necessary to investigate a better way to dispose of molluscs in IMTA using biofloc, towards a reduce stress factors to these organisms.

## CONCLUSIONS

In summary, the concentration of cyanobacteria, especially the genera *Aphanocapsa* sp. and *Pseudanabaena* sp., negatively affected the zootechnical performance of shrimp and molluscs cultured integrated in a BFT system. However, the integrated cultures showed advantages in the shrimp zootechnical performance and presented a lower concentration of cyanobacteria - due to the filtration by the molluscs. Finally, West Indian pointed venus showed low survival in biofloc system regardless of stocking density.

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