

ARTICLE

Antibacterial activity in three *Chaetoceros* microalgae species cultures by using antibiotics

Actividad antibacteriana en cultivos de tres especies de microalgas *Chaetoceros* utilizando antibióticos

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Resumen.- Las diatomeas, como *Chaetoceros*, crecen en una relación mutualista con bacterias. Sin embargo, en algunos casos es necesario que proliferen en cultivos libres de bacterias. Para reducir la carga de bacterias se usan antibióticos y en determinadas ocasiones se requiere utilizar una mezcla con más de un antibiótico. El objetivo de este trabajo fue obtener un protocolo rápido y efectivo para reducir la carga bacteriana y evaluar la respuesta en crecimiento de tres especies de *Chaetoceros* de importancia acuícola. Se utilizaron antibióticos individuales y en mezcla. El crecimiento de microalgas y bacterias heterotróficas fue evaluado. Los parámetros de crecimiento muestran mayor concentración para *C. muelleri* ($3,15 \times 10^6$ céls mL⁻¹) y bajos valores para *C. calcitrans* ($2,98 \times 10^6$ céls mL⁻¹). La mayor tasa de crecimiento fue para *C. calcitrans* (0,77 divisiones por día) y los menores valores fueron para *Chaetoceros* sp. (0,60 divisiones por día). Los parámetros de crecimiento en bacterias heterotróficas muestran una carga bacteriana significativamente mayor para *Chaetoceros* sp. ($19,16 \times 10^6$ UFC (UFC, unidades formadoras de colonias) mL⁻¹) y los menores valores fueron para *C. calcitrans* ($12,23 \times 10^6$ UFC mL⁻¹). La tasa de crecimiento de las bacterias heterotróficas presentes en los cultivos de *Chaetoceros* fue similar entre las tres especies estudiadas. La estreptomycin[®] y el sulfato G41[®] producen reducción parcial de la carga bacteriana. El tratamiento más efectivo para las tres especies fue uso de una mezcla de antibióticos compuesta de ampicilina[®] (250 µg mL⁻¹), kanamicina[®] (200 µg mL⁻¹), neomicina[®] (50 µg mL⁻¹) y estreptomycin[®] (100 µg mL⁻¹) durante tres días. La mezcla preparada con alta concentración de antibióticos produjo una reducción de la carga bacteriana (100%) por tres días, sin embargo, también redujo significativamente el crecimiento (10 a 30%) de las tres especies de *Chaetoceros*.

Palabras clave: Carga bacteriana, *Chaetoceros* spp., antibióticos, cultivos axénicos, bacterias heterotróficas

Abstract.- Diatoms, such as *Chaetoceros*, grow in a mutualistic relationship with bacteria. However, in some cases, it is necessary to grow them in bacteria-free cultures. To reduce bacterial load, antibiotics are used, and on certain occasions it is necessary to use a mixture with more than one antibiotic. This work aimed to obtain a quick and effective protocol to reduce the bacterial load and evaluate the response of three *Chaetoceros* species with aquacultural importance. Single and mix antibiotics were used. Microalgal and bacterial growth was measured. The growth parameters for diatoms showed that the significantly highest cell concentration was for *C. muelleri* (3.15×10^6 cells mL⁻¹) and the lowest values to *C. calcitrans* (2.98×10^6 cells mL⁻¹). The significantly highest growth rate was for *C. calcitrans* (0.77 divisions per day), and the lowest values for *Chaetoceros* sp. (0.60 divisions per day). The growth parameters for heterotrophic bacteria showed that the significantly highest bacterial load was for *Chaetoceros* sp. (19.16×10^6 CFU (Colony-Forming Units) mL⁻¹) and the lowest values were for *C. calcitrans* (12.23×10^6 CFU mL⁻¹). The growth rate of the heterotrophic bacteria present in *Chaetoceros* cultures was similar among the three studied species. Streptomycin[®] and sulfate G41[®] produced a partial reduction of bacterial load. The most effective treatment for all three species was the use of an antibiotic mix composed of ampicillin[®] (250 µg mL⁻¹), kanamycin[®] (200 µg mL⁻¹), neomycin[®] (50 µg mL⁻¹), and streptomycin[®] (100 µg mL⁻¹) for three days. The mix prepared with the highest antibiotic concentration produced a reduction of bacteria (100%) for three days; however, it also induced a significant reduction of the growth of the three *Chaetoceros* species.

Key words: Bacterial load, *Chaetoceros* spp., antibiotics, axenic cultures, heterotrophic bacteria

INTRODUCTION

Diatoms are highly diverse photosynthetic organisms characterized by its silica frustules (Leynaert *et al.* 2018). They are cosmopolite organisms that can be found in marine and seawater environments and are responsible for 20-25% of global primary production (Bozarth *et al.* 2009). Among all known diatoms, *Chaetoceros* is a primarily marine genus and one of the most abundant, reaching almost 228

described species (Guiry & Guiry 2021). Species belonging to this genus, such as *Chaetoceros muelleri* (Kumaran *et al.* 2017), *Chaetoceros calcitrans* (Milagros *et al.* 2018), and *Chaetoceros* sp. (Sánchez-Saavedra & Voltolina 1995, 2006), have a great importance to aquaculture industry due to their biochemical composition, making them an excellent source of feed to fish, crustaceans, and mollusk larvae. The strain of *Chaetoceros* sp. (CHX1) was isolated

in 1988 from Todos Santos Bay at Baja California, Mexico (Trujillo-Valle 1993). This strain has a high potential to be used on aquaculture due its high population growth rates and biochemical composition (Sánchez-Saavedra & Voltolina 1995, 2001, 2006). The species identification has remained uncertain, but it seems to be a strain of *Chaetoceros gracilis* (Tapia-Gallardo 2019).

Diatoms coexist with other microalgal species and bacteria. In fact, exopolysaccharides (EPS) produced by diatoms are utilized by heterotrophic bacteria as their source of organic carbon (Duff *et al.* 1996, Johansson *et al.* 2019, Koedooder *et al.* 2019). Thus, diatom cultures commonly have a high content of heterotrophic bacteria, which may have deleterious effects on growth performance, nutritional quality, and safety of the cells if they are used as feed in aquaculture (Molina-Cárdenas *et al.* 2016). Bacteria may even cause interference in biotechnology, pharmaceutical, biochemical, physiological, taxonomic, and genetic studies, such as the production of compounds with biological activity or genome of diatom species (Vu *et al.* 2018). To reduce the negative impact of high bacterial load it has been recommended to use axenic cultures (*i.e.*, free from undesirable contaminants like bacteria, fungi) (Cho *et al.* 2013). However, the production and maintenance of axenic cultures is often difficult and a time-consuming task (Bruckner & Kroth 2009, Lee *et al.* 2015, Zakharova *et al.* 2020).

Several physical and chemical strategies, involving centrifugation, selective growth media, UV irradiation, filtration, sonication, dilution, purification by micropipette, vortexing, application of mixtures and doses of antibiotics (Andersen & Kawachi 2005, Bruckner & Kroth 2009) are frequently used to control or reduce the bacterial load in microalgal cultures (Sena *et al.* 2011, Molina *et al.* 2019). In some cases, the use of only one technique is not enough to reduce the bacterial load and the application of two or more techniques is required. Additionally, specialized equipment and trained personnel are required (Ishii *et al.* 2018).

Some efforts to obtain axenic cultures of the diatoms *Nitzschia capitellata* and *Halamphora coffeiformis* (Guiry & Guiry 2021) have been performed by pipetting, seeding in agar, use of detergents, phenol solutions, and UV radiation. The results were not satisfactory, and the diatom cultures were still contaminated with bacteria and the microalgae cells were damaged (Jones *et al.* 1973). In a study performed by Nagai *et al.* (1998) the application of washes with a sieve and antibiotic treatment was effective to obtain axenic cultures of the diatom *Coscinodiscus wailesii*. Recently Ishii *et al.* (2018) reported a method to obtain axenic cultures of several microalgal species by sieving and washing resting stage cells. The method was successful to remove bacteria from several diatom species, but the authors affirm that the

protocol is useful to species that have endogenous resting spores and cannot be applied to other diatoms.

Although the use of antibiotics is the most common technique employed to reduce bacterial load (Choi *et al.* 2008, Han *et al.* 2014), effectiveness of bacteria removal from the microalgae cultures can depend on the mechanism of action of the antibiotics used and the bacteria associated to diatom cultures. Thus, the selection is a critical issue when a protocol to reduce bacterial load in diatom cultures has to be decided, in order to remove bacteria and avoid deleterious effects on microalgae (Han *et al.* 2014). Additionally, the response is species-specific and protocols should thus be adapted to each microalgal species.

This work aimed to obtain a quick and effective protocol to reduce bacterial load and evaluate growth response of *Chaetoceros muelleri*, *Chaetoceros calcitrans*, and *Chaetoceros* sp., which are widely used in aquaculture facilities due to their good growth performance, nutritional value and for being organisms of interest for genetic studies.

MATERIALS AND METHODS

DIATOMS CULTURES

Chaetoceros muelleri (CHM1), *Chaetoceros calcitrans* (CHC1) and *Chaetoceros* sp. (CHX1) strains were provided by the microalgal collection of CICESE, Department of Aquaculture. Cultures of each *Chaetoceros* were maintained in batch using triplicate sets in 125 mL Erlenmeyer flasks with 100 mL “P” media (Guillard & Ryther 1962). Light was provided by fluorescent lamps at 50 $\mu\text{mol m}^{-2} \text{s}^{-1}$, 21 ± 1 °C and salinity of 35. To characterize culture growth, cell concentration and heterotrophic bacteria were measured daily.

MICROALGAE COUNTS

The cell concentration of each strain of *Chaetoceros* cultures was measured by direct counts with a hemocytometer (Bright Line, Hausser Scientific, USA). The cell concentrations were \log_2 -transformed to determine the growth rate (μm : divisions per day) of each strain of *Chaetoceros*, according to the following equation, as described by Fogg & Thake (1987).

$$\mu\text{m} = [\text{Log}_2(N_2) - \text{Log}_2(N_1)] / (t_2 - t_1) \quad (1)$$

where, N_1 and N_2 are the cell concentrations at the initial day (t_1) and final day (t_2) respectively, measured during the exponential growth.

BACTERIAL COUNTS

The concentration of heterotrophic bacteria in the cultures of each strain of *Chaetoceros* was obtained daily. The content of bacteria was estimated from the culture medium and dilutions of 10^{-3} were made for each case. Dilutions for bacterial counts were carried out using a physiological serum (9 g of NaCl L⁻¹), and, for each dilution, 0.1 ml was used to inoculate a Petri dish with 20 mL of the ZoBell medium (ZoBell 1941). Petri dishes were incubated in a mini VWR incubator at 31 °C for 48 h. Bacterial load was expressed as colony-forming units per milliliter (CFU mL⁻¹) (Gerhardt *et al.* 1981) and was used to determine the growth rate of bacteria (μ_b) according to equation 1, this growth model was used for bacteria with Log₂ because they have binary division.

BACTERIAL LOAD REDUCTION ASSAYS

The three *Chaetoceros* species were maintained as monospecific batch cultures in 10 mL glass tubes with 5 mL “f” media (Guillard & Ryther 1962). The culture conditions were the same as described for diatom culture.

To decrease bacterial load of the cultures, two assays were carried out. The first assay included 5 treatments (Table 1) to select the best protocol for reducing bacterial load while maintaining microalgal growth. The second assay considered three treatments (Table 1) to see the effect of antibiotics on bacterial load while maintaining microalgal growth. All treatments were applied for 48 h.

The first assay consisted of:

1) Treatment 1: The cells were concentrated by centrifugation at 2,232 g for 10 min at 4 °C and then, cell package of each culture was washed with sterile seawater.

2) Treatment 2: The cells were concentrated and washed as described for treatment 1. The collected cell package was suspended in 5 mL of “f” media with the addition of 75 μ g mL⁻¹ of streptomycin® (SIGMA).

3) Treatment 3: The cells were concentrated and washed as described for treatment 1. The collected cell package was suspended in 5 mL of “f” media with the addition of 250 μ g mL⁻¹ of sulfate G41® (SIGMA).

4) Treatment 4: The cells were concentrated and washed as described for treatment 1. The collected cell package was suspended in 5 mL of “f” media with the addition of

an antibiotic mix composed of ampicillin® (SIGMA) 250 μ g mL⁻¹, kanamycin® (SIGMA) 200 μ g mL⁻¹, neomycin® (SIGMA) 50 μ g mL⁻¹, and streptomycin® (SIGMA) 100 μ g mL⁻¹.

5) Treatment 5: Control of the treatments, 5 mL of each culture was maintained with “f” media in 10 mL glass tubes in triplicates and without the application of any of the four treatments described above.

A second assay was performed based on the previous results obtained from the first assay (Treatment 4), to determine whether a higher antibiotic mix concentration could reduce the bacterial count without damaging the microalgae (Table 1).

A) Treatment A consisted of washing the cell package as described for treatment 1. The collected cell package was suspended in 5 mL of “f” media, with the addition of the antibiotic mix composed of ampicillin® 250 μ g mL⁻¹, kanamycin® 200 μ g mL⁻¹, neomycin® 50 μ g mL⁻¹, and streptomycin® 100 μ g mL⁻¹.

B) Treatment B consisted of washing the cell package as described for treatment 1. The collected cell package was suspended in 5 mL of “f” media with the addition of an antibiotic mix composed of ampicillin® 500 μ g mL⁻¹, kanamycin® 400 μ g mL⁻¹, neomycin® 100 μ g mL⁻¹ and streptomycin® 200 μ g mL⁻¹.

C) Treatment C as control of the treatments, 5 mL of each culture was maintained with “f” media in 10 mL glass tubes in triplicates and without the two treatments described above.

STATISTICAL ANALYSIS

All data were tested for homoscedasticity and normality. To analyze the effects of antibiotics on three *Chaetoceros* species a student’s t-test was applied to initial and final cell densities. A one-way ANOVA was used to evaluate the differences in cell concentration to each *Chaetoceros* strain maintained with different treatments. The same analysis was used to obtain the differences in heterotrophic bacteria content among treatments. When significant differences were detected, Tukey *a posteriori* test was used. The significance level for all the analyses was set to $P < 0.05$. The figures were performed using SigmaPlot 10 software (SigmaPlot 2021)¹.

¹SigmaPlot 2021. SYSTAT Software, San Jose, CA. <www.systatsoftware.com>

Table 1. Treatments used to reduce the bacterial load in cultures of three *Chaetoceros* species / Tratamientos usados para reducir la carga bacteriana en cultivos de tres especies de *Chaetoceros*

	Wash and centrifugation	Antibiotic	Concentration ($\mu\text{g mL}^{-1}$)
Assay 1			
1	2232 g for 10 min	None	None
2	2232 g for 10 min	Streptomycin®	75
3	2232 g for 10 min	Sulfate G41®	250
4	2232 g for 10 min	Ampicillin®	250
		Kanamycin®	200
		Neomycin®	50
		Streptomycin®	100
5	None	None	None
Assay 2			
A	2232 g for 10 min	Ampicillin®	250
		Kanamycin®	200
		Neomycin®	50
		Streptomycin®	100
B	2232 for 10 min	Ampicillin®	500
		Kanamycin®	400
		Neomycin®	100
		Streptomycin®	200
C	None	None	None

RESULTS

MICROALGAE AND BACTERIAL GROWTH

The growth curve shows that *C. muelleri* and *C. calcitrans* had an exponential growth phase until day 4, after which the stationary-growth phase began, and during day 7, a decrease in cell concentration was evaluated. *Chaetoceros* sp. maintain their exponential growth from day 2 to day 7 (Fig. 1).

The growth parameters for diatoms show that the significantly highest cell concentration ($P < 0.05$) was observed for *C. muelleri* (3.15×10^6 cells mL^{-1}) and the lowest values for *C. calcitrans* (2.98×10^6 cells mL^{-1} ; Table 1). The significantly highest growth rate was for *C. calcitrans* (0.77 divisions per day) and the lowest values

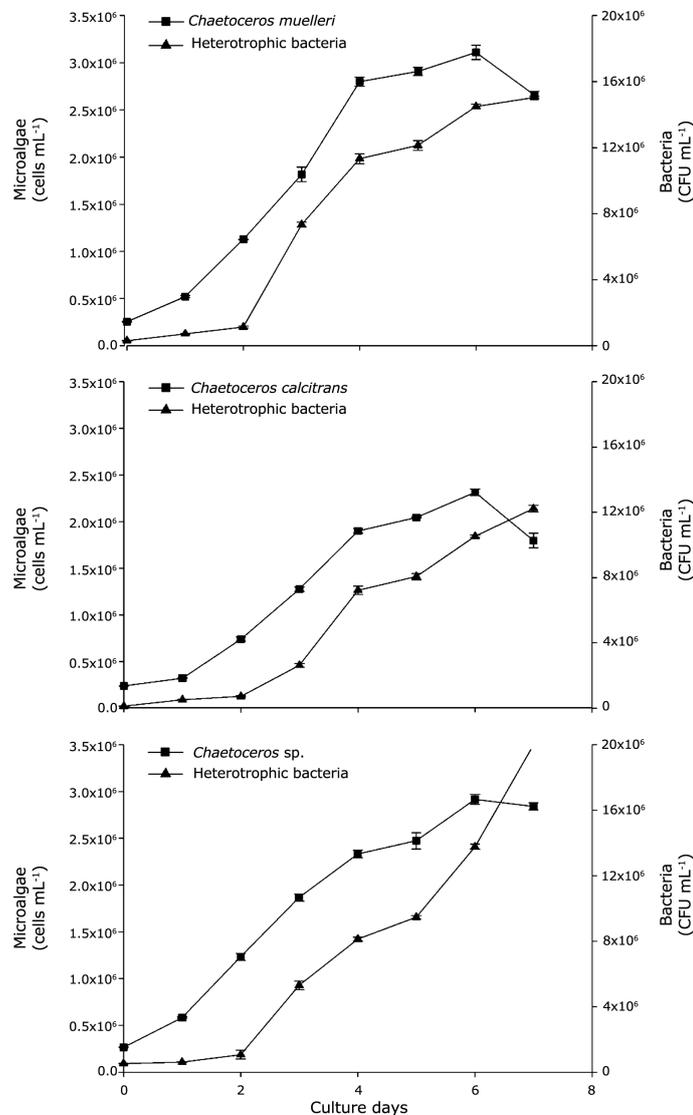


Figure 1. Mean values and standard deviation of cell concentration and heterotrophic bacterial load of monospecific and non-axenic batch cultures of *Chaetoceros muelleri*, *C. calcitrans* and *Chaetoceros* sp. / Valores promedio y desviación estándar de la concentración de células y carga de bacterias heterotróficas de cultivos en lote mono-específicos, no axénicos de *Chaetoceros muelleri*, *C. calcitrans* y *Chaetoceros* sp.

for *Chaetoceros* sp. (0.60 divisions per day; $P < 0.05$). The growth parameters for heterotrophic bacteria show that the significantly highest bacterial load was for *Chaetoceros* sp. (19.16×10^6 CFU mL⁻¹) and the lowest values for *C. calcitrans* (12.23×10^6 CFU mL⁻¹; $P < 0.05$). The growth rate of the heterotrophic bacteria was similar between the three *Chaetoceros* species ($P > 0.05$; Table 2).

BACTERIAL LOAD REDUCTION ASSAYS

In the first assay, for *C. muelleri* cultures maintained with the different treatments to decrease bacterial load, the results showed that the final cell concentration (FCC) was significantly higher when washed with sterile water + sulfate G41® (T3: 3.36×10^6 cells mL⁻¹). Meanwhile, the lowest cell concentration was for the control without antibiotic treatment (control) (T5: 2.43×10^6 cells mL⁻¹; $P < 0.05$; Table 3). The treatment that significantly reduced

the heterotrophic bacterial load corresponded to washing with sterile seawater + antibiotic mix (T4: 0.08×10^6 CFU mL⁻¹), meanwhile, the highest values were for the control treatment (T5: 11.74×10^6 CFU mL⁻¹; $P < 0.05$; Table 3).

For *C. calcitrans* cultures, the final cell concentration (FCC) was significantly higher for washing with sterile seawater + streptomycin® (T2) and washing with sterile seawater + antibiotic mix (T4) (both with 2.78×10^6 cells mL⁻¹) and washed with sterile seawater + sulfate G41® (T3: 2.86×10^6 cells mL⁻¹); meanwhile the lowest cell concentration was for the control without antibiotic treatment (control) (T5: 2.33×10^6 cells mL⁻¹; $P < 0.05$; Table 3). The treatment that significantly reduced the heterotrophic bacterial load corresponded to washing with sterile seawater + antibiotic mix (T4: 0.06×10^6 CFU mL⁻¹), meanwhile, the highest values were for the control treatment (T5: 8.60×10^6 CFU mL⁻¹) ($P < 0.05$) (Table 3).

Table 2. Mean values and standard deviation of maximum cell concentration (MCC x 10⁶ cells mL⁻¹), and heterotrophic bacteria load (HBL x 10⁶ CFU mL⁻¹), growth rate of microalgae (μm, divisions per day) and growth rate of heterotrophic bacteria (μb, divisions per day) of *Chaetoceros muelleri*, *C. calcitrans* and *Chaetoceros* sp. of monospecific batch cultures (first assay) / Valores promedio y desviación estándar de la concentración máxima de células (MCC x 10⁶ céls mL⁻¹) y carga de bacterias heterotróficas (HBL x 10⁶ UFC mL⁻¹), tasa de crecimiento de microalgas (μm, divisiones por día) y bacterias heterotróficas (μb, divisiones por día) de *Chaetoceros muelleri*, *C. calcitrans* y *Chaetoceros* sp. de cultivos monoespecíficos en lotes (primer ensayo)

Species	Growth parameters			
	Microalgae		Bacteria	
	MCC	μm	HBL	μb
<i>C. muelleri</i>	3.15 ± 0.04 a	0.73 ± 0.01 b	16.15 ± 0.13 b	0.95 ± 0.01 a
<i>C. calcitrans</i>	2.34 ± 0.01 c	0.77 ± 0.01 a	12.23 ± 0.20 c	0.98 ± 0.04 a
<i>Chaetoceros</i> sp.	2.98 ± 0.05 b	0.60 ± 0.01 c	19.96 ± 0.15 a	0.93 ± 0.02 a

Different letters denote significant differences (one-way ANOVA $P < 0.05$: a>b>c)

Table 3. Mean values and standard deviation of initial cell concentration (ICC x 10⁶ cells mL⁻¹), final cell concentration (FCC x 10⁶ cells mL⁻¹) and heterotrophic bacteria load (HBL x 10⁶ CFU mL⁻¹) of *Chaetoceros muelleri*, *C. calcitrans* and *Chaetoceros* sp. of monospecific batch cultures with different treatments (first assay) / Valores promedio y desviación estándar de la concentración celular inicial (ICC x 10⁶ céls mL⁻¹), concentración celular final (FCC x 10⁶ céls mL⁻¹) y carga de bacterias heterotróficas (HBL x 10⁶ UFC mL⁻¹) de *Chaetoceros muelleri*, *C. calcitrans* y *Chaetoceros* sp.) de cultivos monoespecíficos en lotes con diferentes tratamientos (primer ensayo)

Species	T1	T2	T3	T4	T5
<i>C. muelleri</i>					
ICC	2.21 ± 0.04 a	2.17 ± 0.07 a	2.08 ± 0.05 a	2.07 ± 0.07 a	2.35 ± 0.17 a
FCC	2.57 ± 0.67 b	3.06 ± 0.08 a	3.36 ± 0.09 a	2.77 ± 0.03 a	2.43 ± 0.02 b
HBL	5.58 ± 0.37 b	11.40 ± 0.56 a	9.60 ± 0.28 ab	0.08 ± 0.02 c	11.74 ± 0.48 a
<i>C. calcitrans</i>					
ICC	2.03 ± 0.08 a	1.94 ± 0.08 a	2.01 ± 0.01 a	1.98 ± 0.01 a	2.11 ± 0.01 a
FCC	2.41 ± 0.01 b	2.78 ± 0.05 a	2.86 ± 0.01 a	2.78 ± 0.05 a	2.33 ± 0.01 b
HBL	4.40 ± 0.14 b	7.10 ± 0.14 a	5.70 ± 0.14 ab	0.06 ± 0.00 c	8.60 ± 0.14 a
<i>Chaetoceros</i> sp.					
ICC	2.05 ± 0.07 a	2.07 ± 0.07 a	2.10 ± 0.03 a	2.11 ± 0.01 a	2.17 ± 0.07 a
FCC	2.61 ± 0.01 b	2.81 ± 0.01 a	2.68 ± 0.01 a	2.91 ± 0.02 a	2.45 ± 0.03 b
HBL	7.57 ± 0.10 b	13.26 ± 0.09 a	11.06 ± 0.09 ab	0.13 ± 0.01 c	15.30 ± 0.14 a

T1, washed with sterile seawater. T2, washed with sterile seawater + streptomycin®. T3, washed with sterile seawater + sulfate G41®. T4, washed with sterile seawater + antibiotic mix. T5, control without treatment addition. Different letters denote significant differences (one-way ANOVA $P < 0.05$: a>b>c)

For *Chaetoceros* sp. cultures, FCC was significantly higher for washing with sterile seawater + antibiotic mix (T4: 2.91×10^6 cells mL⁻¹) and washing with sterile seawater + antibiotic mix (T4: 2.86×10^6 cells mL⁻¹); meanwhile, the lowest cell concentration was for the control without antibiotic treatment (control) (T5: 2.45×10^6 cells mL⁻¹; $P < 0.05$; Table 3). The treatment that significantly reduced the heterotrophic bacteria load was for washing with sterile seawater + antibiotic mix (T4: 0.13×10^6 CFU mL⁻¹); meanwhile, the highest values were for the control treatment (T5: 15.30×10^6 CFU mL⁻¹; $P < 0.05$; Table 3).

For the second assay, the antibiotic mix used to decrease the bacterial load on the diatom cultures showed that the significantly highest cell concentration corresponded to control treatment of the three strains ($P < 0.05$; Fig. 2). The control treatment increased the cell concentration throughout the culture time until day 5, after which the cell concentration decreased in the three diatom strains. The heterotrophic bacterial load increased significantly ($P < 0.05$) throughout the culture time on the control treatment.

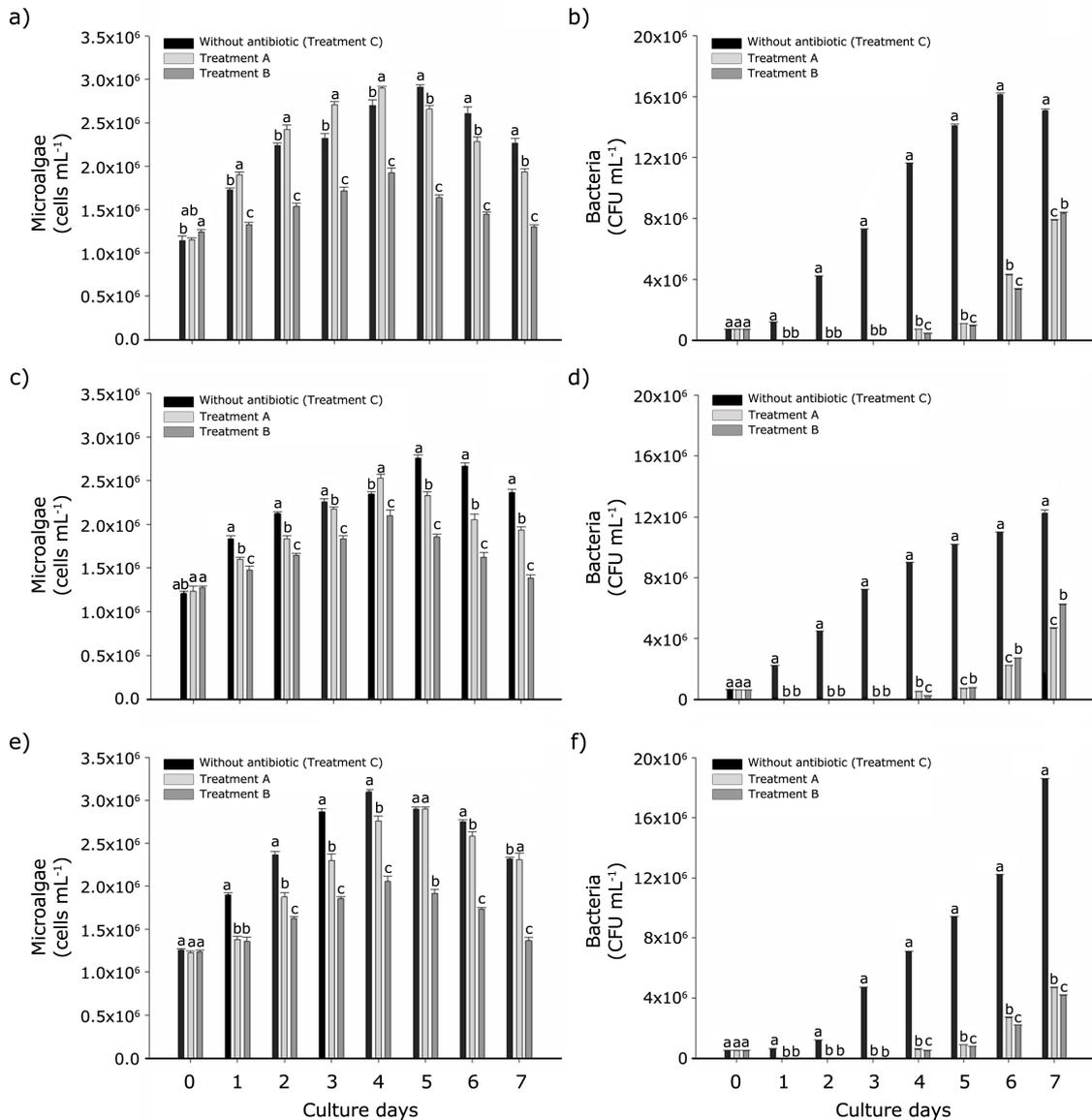


Figure 2. Mean values and standard deviation of cell concentration and heterotrophic bacteria load of monospecific batch cultures of *Chaetoceros muelleri* (a, b), *C. calcitrans* (c, d) and *Chaetoceros* sp. (e, f) maintained with three treatments (second assay). Treatment A: ampicillin® 250 µg mL⁻¹, kanamycin® 200 µg mL⁻¹, neomycin® 50 µg mL⁻¹, and streptomycin® 100 µg mL⁻¹. Treatment B: ampicillin® 500 µg mL⁻¹, kanamycin® 400 µg mL⁻¹, neomycin® 100 µg mL⁻¹, and streptomycin® 200 µg mL⁻¹. Treatment C: culture without antibiotic addition (second assay). Different letters denote significant differences (one-way ANOVA $P < 0.05$: $a > b > c$) / Valores promedio y desviación estándar de la concentración de células de *Chaetoceros muelleri* (a, b), *C. calcitrans* (c, d) y *Chaetoceros* sp. (e, f), y bacterias heterotróficas mantenidas con dos tratamientos. Tratamiento A: ampicilina® 250 µg mL⁻¹, kanamicina® 200 µg mL⁻¹, neomicina® 50 µg mL⁻¹ y estreptomicina® 100 µg mL⁻¹. Tratamiento B: ampicilina® 500 µg mL⁻¹, kanamicina® 400 µg mL⁻¹, neomicina® 100 µg mL⁻¹ y estreptomicina® 200 µg mL⁻¹. Tratamiento C: cultivo sin adición de antibiótico (segundo ensayo). Letras diferentes denotan diferencias significativas (ANOVA de una vía $P < 0,05$: $a > b > c$)

In the second assay, the lowest concentration mixture (treatment A) slightly diminished the cell concentration of the three *Chaetoceros* strain with respect to control treatment (Fig. 2). The heterotrophic bacterial load decreased significantly ($P < 0.05$) on the three diatom strains until day 3, and then, slowly increased until day 7.

The antibiotic mix with the highest concentration showed a significant decrease in cell concentration (10 to 30%) for the three diatom strains throughout the culture time ($P < 0.05$) (Fig. 2). The heterotrophic bacterial load significantly decreased ($P < 0.05$) on the three diatom strains until day 3; after that, it slowly increased until day 7 with similar or lower bacterial load as described with the low concentration of the antibiotic mix.

DISCUSSION

MICROALGAE AND BACTERIAL GROWTH

Microalgae cultures contain bacteria that interact with each other, associated with mechanisms that range from the production of compounds that inhibit or stimulate the growth of microalgae species. This type of interspecific relationship between species of bacteria-microalgae has been seen to depend on mutualism in many cases (Hernández-Pérez & Labbé 2014, Clarke *et al.* 2019). The control of both bacterial and microalgal growth affects the growth, composition, and production of metabolites, and can result in great biotechnological benefits (Réveillon *et al.* 2016, Koedoeder *et al.* 2019).

The cell concentration and growth rate measured on the cultures of *C. muelleri* were lower than those obtained by other authors (López-Eliás *et al.* 2005, Pacheco-Vega & Sánchez-Saavedra 2009, Orozco-Borbón *et al.* 2014), and the differences were due to the use of high culture volume, high temperature, and high cell concentration used as inoculum to the cultures. For *C. calcitrans* the growth rate was higher than the one obtained by Phatarpekar *et al.* (2000); meanwhile, cell concentration was lower than measured by Villa *et al.* (2014). For *Chaetoceros* sp., the cell concentration was lower than the concentration obtained by Sánchez-Saavedra & Voltolina (2006) when the cultures were maintained with irradiance 60% higher than the irradiance used on this work.

The content of heterotrophic bacteria of the three *Chaetoceros* species used in this work increased proportionally as the cell concentration of the diatoms augmented. In *C. muelleri* cultures it has been described that the growth of heterotrophic bacteria rises directly proportionally as an increase in the cell concentration of the diatom throughout the culture time (Orozco-Borbon *et al.* 2014). These trends were previously described by other authors for different microalgae strains (Gomez-Gil

et al. 2002, Ruiz-Güereca & Sánchez-Saavedra 2016). For *C. calcitrans*, the heterotrophic bacteria content was previously described as 10^4 to 10^7 CFU mL⁻¹ in cultures maintained in a commercial shrimp hatchery (Lizárraga-Partida *et al.* 1997).

In some cases, non-pathogenic bacteria can increase the growth of microalgae cultures such *Nannochloropsis oculata*, and *C. calcitrans*, which significantly increase their growth, when they are maintained in co-culture with *Bacillus* sp. and *Pseudomonas* sp. obtained from algae culture system (Sureshkumar *et al.* 2014). However, the impact of algal growth rates on the bacterial load in six microalgae species, shows that higher bacterial levels were associated with slow-growing microalgae on a per volume basis. The microalgae cultures can reach the contents of bacterial loads as 1 to 10^8 CFU per liter of culture (Murchelano & Brown 1969). Variations on the bacterial load were more related to species. The higher contents of bacterial loads were associated with the Bacillariophyceae group reaching values between 5.7 to 16.4 CFU algal cell⁻¹, and for other microalgae species, the bacteria load was 0.2 to 4.3 CFU algal cell⁻¹ (Salvesen *et al.* 2000).

Bacterial growth rate was similar in all three *Chaetoceros* strains cultures. However, the density of the heterotrophic bacteria was higher with *Chaetoceros* sp., followed by the values measured with *C. calcitrans*, and *C. muelleri*. This difference in bacterial content between the three *Chaetoceros* species can be attributed to a species-specific interaction due to the specific extracellular products produced by each diatom species. In some cases, the bacterial content associated with different microalgae strains is related to the production of antibacterial substances, which are generally most abundant in slow-growing microalgae cultures and the early stationary growth phase when the competition for nutrients is high due to abundance of both microalgae and bacteria (Borowitzka 1995). However, studies are needed to determine the chemical characterization of the compounds produced by each one of the *Chaetoceros* species and their associated bacteria.

The importance of maintaining monospecific microalgal cultures with low bacterial load is that the bacterial content can modify the growth characteristics and biochemical composition of the microalgal cells. Additionally, it is important to maintain cultures with low bacterial content or axenic cultures for studies such as genome sequencing (Vu *et al.* 2018, Pinder *et al.* 2019), interaction between species, harmful algal blooms, microalgae-bacteria consortia, cell cryopreservation, identification of bioactive compounds for biotechnology or pharmacology (Guillard 2005, Borowitzka 2013, Vu *et al.* 2018).

BACTERIAL LOAD REDUCTION ASSAYS

The goal of purification methods is to obtain a viable culture of a single microalgal species, free of all other species (Guillard 2005).

In the first assay of this study, a lower content of heterotrophic bacteria in the monospecific cultures of *Chaetoceros* was obtained from the application of the antibiotic mix (treatment 4: ampicillin® 250 µg mL⁻¹, kanamycin® 200 µg mL⁻¹, neomycin® 50 µg mL⁻¹ and streptomycin® 100 µg mL⁻¹). This result is due to the interaction between the antibiotics used and the mechanism of action of each antibiotic, resulting in low concentration of bacteria and no apparent damage to the microalgae cells (Molina-Cárdenas *et al.* 2016).

The use of centrifugal washes, as well as the application of a single antibiotic, has been reported to have a low efficiency to achieve axenic cultures of several microalgae species (Wilkens & Maas 2012). The disadvantage of using centrifugal washes is that the bacterial load of microalgae cultures is reduced for a very short time. It is mentioned that the use of different antibiotics must be specific to each microalgal species, depending on the characteristic bacterial community associated with the microalgae (Bruckner & Kroth 2009). Centrifugation is perhaps a more frequently used method to concentrate the target organism rather than establishing unialgal cultures and avoiding bacteria contamination (Andersen & Kawachi 2005, Molina *et al.* 2019).

According to the results obtained with three *Chaetoceros* species and with the use of the antibiotic mix in two different concentrations, a decrease in the content of heterotrophic bacteria (at levels not detectable for 4 days) was found. However, a reduction of the three *Chaetoceros* species growth (Fig. 2) was observed with the highest antibiotic concentration (treatment B).

The use of antibiotics must be strict and careful because it can increase the resistance in the populations of some bacterial species. Besides, it can affect the cell membrane and thereby inhibit microalgal growth, resulting in axenic cultures with average cell densities of 10⁶ cells mL⁻¹ (González-Pleiter *et al.* 2013, Molina-Cárdenas *et al.* 2016). The use of a mixture of antibiotics as a control of bacteria in microalgae cultures shows synergistic effects that usually increase their toxicity to bacteria. However, some mixtures of antibiotics can produce a decrease in the microalgae population (González-Pleiter *et al.* 2013) or their concentrations, as observed in this study (second assay and treatment B).

Chlorotetracyclines and oxytetracyclines belong to tetracyclines, and are used as antibiotics for controlling Gram-positive and Gram-negative bacteria, mycoplasma, and viruses in the pig farming industry (Arsenakis

2018). These antibiotics act on the inhibition of protein synthesis and prevent the association of aminoacyl-t RNA and the ribosome of bacteria. Enrofloxacin is a broad-spectrum quinolone that is widely used for the control of microorganisms (Guo & Chen 2002). In microalgae, the main approach that has been given to studies related to the use of antibiotics is associated with β-lactams (ampicillin penicillin, chlortetracycline, and oxytetracycline), quinolones (norfloxacin, ciprofloxacin, and enrofloxacin) and macrolides (erythromycin). Our results demonstrate that the use of some antibiotics provided individually or in a mix, can decrease the bacterial load. Interestingly, the antibiotics significantly increase the cell concentration for *C. muelleri* cultures on days one to four. The positive effect of the mix of antibiotics to *C. muelleri* can be due to the cellular lysis of bacteria and that the cell components could be used by the diatom cells as nutrients, thus inducing cell production. The strains of *C. calcitrans* and *Chaetoceros* sp. do not show a growth-promoting effect due to the antibiotic mix. In some cases, the microalgae cells are resistant to some antibiotics at a determined concentration, period of exposure, culture conditions that can inhibit the effect of antibiotics and tolerance of some species of microalgae due to the physiological characteristics of the cells (Qunying & Aiyi 2000, Guillard 2005). Penicillin could inhibit the growth of *Phaeodactylum tricorutum* and the inhibition decreased after five days. However, the use of penicillin increased the growth of *Isochrysis* sp., *Isochrysis galbana* and *Platymonas subcordiformis*. The use of streptomycin can inhibit the growth of *Phaeodactylum tricorutum*, *Isochrysis* sp., *Isochrysis galbana*, and *Platymonas subcordiformis*. The use of chloramphenicol inhibits the growth of *Isochrysis* sp. and *Isochrysis galbana* (Qunying & Aiyi 2000). Some antibiotics act as growth promoters in agriculture due to the effect of lack of competition related to nutrients (Dibner & Richards 2005).

The lethality of the antibiotic treatment in microalgae cultures is an intensity-time relationship, the intensity being the dose (concentration) of antibiotics and the time being the period of exposure before transfer to antibiotic-free medium. The fundamental choice is which antibiotic to employ, at what concentration, and for how long. The action of a mixture of antibiotics is not, in general, the sum of individual actions (Guillard 2005). It was found that the three *Chaetoceros* species used in this work have different communities of bacteria, due to the different bacterial load throughout the culture time and to differences in susceptibility to decrease bacterial load or induce the growth of some diatom species. It has been found that the use of antibiotics to control bacterial growth in microalgal cultures - provided individually, in a binary mixture, or a mix - produces an inhibitory effect of various species of bacteria. However, the use of antibiotics is more effective in mixtures. In fact, there is a relationship between

concentration, duration of treatment, and environmental factors, such as temperature and light (Carusso *et al.* 2018, Rico *et al.* 2018).

The importance of carrying out studies of axenization of microalgae species is, among others, to be used in genetic, transcriptomic, and microbiome studies, as well as in the study of the interrelation of bacteria and microalgae (Tapia *et al.* 2016, Severin & Erdner 2019, Burgunter-Delamare *et al.* 2020).

In conclusion, it was found that the use of streptomycin produces bacterial load reduction on the three *Chaetoceros* species: 3% for *C. muelleri*, 17.5% for *C. calcitrans*, and 14% for *Chaetoceros* sp. When washing with sterile seawater was used, heterotrophic bacteria decreased nearly 50% in the three *Chaetoceros* cultures. The use of sulfate G41® produces a reduction of bacterial load of 18% for *C. muelleri*, 33% for *C. calcitrans*, and 24% for *Chaetoceros* sp. The most effective treatment was the use of an antibiotic mix composed of ampicillin® (250 µg mL⁻¹), kanamycin® (200 µg mL⁻¹), neomycin® (50 µg mL⁻¹), and streptomycin® (100 µg mL⁻¹), because this antibiotic mixture induces a reduction of bacterial load (100%) in the three *Chaetoceros* species from day one to day three. The growth rate of the diatoms was slightly reduced with respect to the control treatment. The mix prepared with the high antibiotic concentration produced a reduction of bacterial load (100%) for three days; however, it also generated a significant reduction of *Chaetoceros* growth (10 to 30%).

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