



Effects of *Bacillus subtilis* C-3102-supplemented diet on growth, non-specific immunity, intestinal morphometry and resistance of hybrid juvenile *Pseudoplatystoma* sp. challenged with *Aeromonas hydrophila*

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Abstract

The aim of the present study was to investigate the influence of dietary supplementation with *Bacillus subtilis* C-3102 on the productive, intestinal histomorphometry, haemato-immunological aspects, as well as evaluated the resistance of juvenile *Pseudoplatystoma* sp. after challenge with *Aeromonas hydrophila*. Inclusion levels of the commercial probiotic were set at 0 (control), 1, 2, 3 and 4% of CALSPORIN® kg feed⁻¹ with six replicates. Blood samples were collected on day 0, 10 and 20 for hematological analysis, and on the 20th day, samples were collected for intestinal histomorphometry, zootechnical and survival analyses. The results showed that probiotic supplementation with *B. subtilis* significantly improved intestinal morphology in fish from groups 1% and 2% and phagocytic activity in all supplemented groups, regardless of the applied dose. In addition, on the 20th day, improvements in non-specific immunity were observed, such as an increase in the number of monocytes (groups 2, 3 and 4%), eosinophils (group 3%), leukocytes (1%) and lymphocytes (group 4%) of the fish that received the supplemented feed. It was also possible to verify that the probiotic caused significant changes in the haemogram between the sampling periods on 0, 10th and 20th day, showing activation of specialized cells of the fish's non-specific defense mechanism. At the end of the experimental period, supplementation of *B. subtilis* improved the productive indexes and survival of *Pseudoplatystoma* sp. after bacterial challenge.

Highlights

- Short term of probiotic feeding evaluation in the diet of Brazilian native catfish *Pseudoplatystoma* sp.
- *B. subtilis* causes significant improvements in non-specific fish immunity.
- Probiotic feeding increases resistance to *A. hydrophila*.
- Short-term supplementation leads to productive improvements in *Pseudoplatystoma* sp.

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Introduction

Environmental issues are increasingly present in the aquaculture production sector, where gaps related to environmental sustainability need answers from research and development groups, as they are extremely important for the sustainable advancement of the activity (Kavitha et al. 2018; Di et al. 2019; Kuebutornye et al. 2019). In this context, probiotics are emerging as environmentally friendly tools for application in aquaculture (Di et al. 2019; Kuebutornye et al. 2019). As a functional additive, probiotics directly affect the performance of aquatic organisms and have been widely used to supplement fish and shrimp diets (Liu et al. 2012; Telli et al. 2014; Kavitha et al. 2018; Di et al. 2019; Kuebutornye et al. 2019; Interaminense et al. 2019; Zhou et al. 2019).

Freshwater fish farming is the activity that most contributes to fish production in the world (FAO 2018). In this scenario, we can highlight the catfish of the genus *Pseudoplatystoma* sp. of high economic value, which emerge as an object of interest for aquaculture in Latin America (Campos 2005; Inoue et al. 2009). However, disease outbreaks, followed by high mortality rates and consequent economic losses, are common and are generally related to nutritional and health factors in the early stages of commercial production of *Pseudoplatystoma* sp. (Tavares-Dias and Martins 2017; Tavares et al. 2018). Nutrition is closely linked to the health of aquatic organisms. Thus, probiotics may, by competing for adhesion of the adhesion site in the digestive tract, nutrients and energy sources, prevent the establishment of pathogens in the intestinal epithelium, prevent injury, restore intestinal mucosa integrity and thereby improve the absorption of nutrients (Tournut 1998; Petri 2000; Kiron 2012; Kuebutornye et al. 2019) and thus may restrict the growth of undesirable microorganisms (Pal et al. 2019).

Considering all the possible nutritional and health benefits, several components have been tested to act on the intestinal mucosa aiming to manipulate the intestinal flora and morphology, to hinder the adhesion and colonization of pathogenic microorganisms to the intestinal epithelium (Tournut 1998; Boleli et al. 2000; Maiorka et al. 2002; Kuebutornye et al. 2019; Owatari et al. 2019). The benefits already observed for different fish species after ingestion of probiotics containing *Bacillus* spp. are diverse, among which we can highlight the increased immune response (Aly et al. 2008; Kumar et al. 2008; Dias et al. 2012; Farias et al. 2016; Nakandakare et al. 2018), growth improvement (Gatesoupe 1999; Farias et al. 2016; Kuebutornye et al. 2019), higher survival against some pathogens (Dias et al. 2012), microbiota replacement after antibiotic use and production of broad-spectrum antimicrobial substances (Green et al. 1999; Raida et al. 2003; Balcázar et al. 2006; Kuebutornye et al. 2019).

Bacteria of the genus *Bacillus* constitute a diverse group of gram-positive rods, characterized by their ability to produce important secondary metabolites such as antibiotics, bioinsecticides, biocins, organic acids and enzymes (Olmos 2003; Silva et al. 2013; Kuebutornye et al. 2019), which are very effective in breaking down a wide variety of carbohydrates, lipids and proteins (Sonnenschein et al. 1993). *Bacillus* are characterized by the formation of resistant spores that allow greater viability after feed processing (Yılmaz et al. 2020). In addition, *Bacillus* has high survival rates after exposure to stomach gastric acid, making them highly usable in aquaculture (Hong et al. 2005; Kuebutornye et al. 2019).

Preliminary studies exposed by Veiga et al. (2020) in a short communication demonstrated some potential effects of *B. subtilis* on juvenile *Pseudoplatystoma* sp., and based on the

information, this study aimed to continue investigations on the influence of *B. subtilis* in the diet of *Pseudoplatystoma* sp., to strengthen the knowledge of its effects on non-specific immunity, haematology, productive performance, intestinal histomorphometry and resistance to *Aeromonas hydrophila*.

Material and methods

The fish used in this study hybrid surubim (*Pseudoplatystoma reticulatum* × *P. corruscans*) were obtained from Pirai Fish Farm, Terenos, MS, Central-Western Brazil. All procedures for fish handling and euthanasia were approved by the Ethical Committee on the Animal Use CEUA/UEMS no 014/2013, using eugenol (75 mg L⁻¹) as an anesthetic procedure.

Experimental design

Water quality monitoring was performed daily at 08:00 with the aid of a handheld YSI Professional Plus multiparameter equipment. The environmental conditions during the experiment remained at a temperature of 24.8 ± 1.8 °C, dissolved oxygen 4.9 ± 0.4 mg L⁻¹, pH 6.9 ± 0.3 and conductivity 193.0 ± 15.0 µS cm⁻¹, within the recommended comfort range for the species (Sipaúba-Tavares et al. 1995).

Three hundred juveniles surubim, trained to receive dry food, with total length and weight of 15.9 ± 0.8 cm and 24.5 ± 3.6 g, respectively, were used. The fish were distributed in 30 experimental units with 40 L volume (10 fish per unit) with continuous water flow and aeration. Before starting the experiment, an external visual check was performed on the fish's body, to verify the absence of lesions or changes in the pigmentation on the skin and gills. The fish remained for 10 days in the acclimatization period, until they got used to the daily handling and feeding routine.

The experimental design was randomized with four treatments and a probiotic-free control group, each with six replications. Four increasing levels of probiotic composed by *B. subtilis* were added to the experimental diet: 1%, 2%, 3% and 4% CALSPORIN® kg feed⁻¹. The probiotic was included in the diet using oily medium (2% soybean oil) according to Nakandakare et al. (2013), for better adherence of the product to feed pellets. An extruded commercial feed 40% crude protein (2–4 mm pellets) was used for carnivorous fish. The fish were fed twice a day until satiety at 7:00 am and 5:00 pm during the experimental period of 20 days. After the inclusion of the probiotic, a sample of the feed from each treatment was submitted to microbiological analysis at the Microbiology Laboratory “Comércio e Indústria Uniquímica Ltda”–Diadema, São Paulo, to quantify the *B. subtilis* concentration present in the feed. For this, a sample of 1 g of feed from each treatment was macerated and inserted in a Falcon tube containing 50 mL of diluent to be homogenized. Subsequently, 1 mL of the sample was serially diluted in Falcon tubes containing 9 mL of diluent solution until the concentration of the dilution reached a count of 30–300 microorganisms per 1 mL. After the final dilution, an aliquot with a volume of 1 mL was sown in a Petri dish containing specific culture medium and incubated in a bacteriological incubator at 37 °C for 48 h. After incubation, *B. subtilis* colonies were confirmed and counted (Table 1).

Animal management and growth performance

To evaluate the growth performance of the fish, biometrics were performed to obtain weight and length data at the beginning and end of the experimental period. The zootechnical indices

Table 1 Microbiological analysis of experimental diets offered to hybrid surubim (*Pseudoplatystoma reticulatum* × *P. corruscans*) supplemented with the probiotic *Bacillus subtilis*

Treatments	CFU <i>B. subtilis</i> g feed ⁻¹	Log ₁₀
Control	ND	ND
1%	8.40 × 10 ⁶	6.9262
2%	1.70 × 10 ⁷	7.2240
3%	3.30 × 10 ⁷	7.5218
4%	3.50 × 10 ⁷	7.5378

CFU colony forming unit, ND not detectable

evaluated were apparent feed conversion (AFC) = feed intake/weight gain; weight gain (WG) = final weight – initial weight; biomass gain (BG) = final biomass – initial biomass; protein efficiency ratio (PER) = ratio between the biomass gain and crude protein intake (CPI), where CPBI corresponds to the percentage of crude protein (CP) of the experimental feed multiplied by the total feed intake; feed intake (FI) = average feed intake per fish during the experimental period; specific growth rate (SGR) = natural log of final weight (g) – natural log of initial weight (g) / experimental period (days) × 100; and survival = the final number of fish / the initial number of fish × 100.

Haematological parameters

For haematological analysis, the blood was collected and analysed at three different times: at the beginning of the experiment (0 day—basal collection), on the 10th day, and on the 20th day. Blood samples were withdrawn from 12 fish per treatment by puncturing the caudal vessel using 3% EDTA emulsified disposable syringes and needles.

The haematological parameters evaluated were hematocrit percentage by the microhematocrit method according to Goldenfarb et al. (1971), hemoglobin according to Collier (1944), red blood cells count (RBC) in a Neubauer chamber. From hematocrit, hemoglobin and erythrocyte amount data, the mean corpuscular volume (MCV) and the mean corpuscular hemoglobin concentration (MCHC) were calculated. Duplicate blood smears were made for each fish and stained with May Grünwald-Giemsa-Wright (Tavares-Dias and Moraes 2003) for differential leukocyte count, total thrombocyte count and white blood cell count (WBC) (Hrube and Smith 1998).

Phagocytic activity

At the end of the probiotic supplementation period, six fish from each experimental group were anesthetized in a solution containing eugenol and then injected intraperitoneally with the *Saccharomyces cerevisiae* yeast solution (9000 cell mm⁻³) according to Ranzani-Paiva et al. (2008).

After 2 h of yeast incubation, the animals were euthanized. Then, a ventral incision was made, where the peritoneal cavity was washed with 1.5 mL of phosphate-buffered saline solution (PBS—137 mM NaCl, 2.7 mM KCl, 20 mM Na₂HPO₄, 5 mM KH₂PO₄, pH 7.4). The fluid containing the inflammatory cells (macrophages) was aspirated with Pasteur pipette and then centrifuged at 1500g for 5 min. After centrifugation, the supernatant was discarded, and the remainder of the precipitate was deposited between the slide and coverslip and observed under phase-contrast microscopy for phagocyte count.

Phagocytic capacity (PC) and phagocytic index (PI) of phagocytes were calculated according to the methodology of Silva et al. (2002, 2005), where phagocytic capacity (PC) is the percentage of a particular cell type that is phagocytizing, expressed by the equation $PC = (\text{number of phagocytes} / \text{total counted phagocytes}) \times 100$, and phagocytic index (PI) is the average number of yeast within phagocytic cells where 100 cells are counted, whose formula is $PI = \text{total number of yeast within phagocytes} / \text{total number of active phagocytes}$.

Intestinal histomorphometry

On the 20th day, biological material was collected for intestinal analysis. Histological slides were made for subsequent histomorphometry analysis of the intestinal epithelium of fish at the Veterinary Pathology Laboratory, UFMS. Twelve animals from each treatment were used.

The fish were euthanized, dissected and submitted to a ventral longitudinal incision for organ exposure. Approximately 3-cm sections were obtained from the midgut, cut longitudinally with microsurgical scissors. Afterward, the samples were fixed open with the internal organ wall facing upwards on a Styrofoam holder, dipped in Bouin's fixative solution for 6 h and then transferred to 70% alcohol. Then, the samples were dehydrated in increasing series of alcohol, diaphanized in xylol and included in paraffin, to obtain cross-sections with the aid of microtome in 5 μm thickness. From each intestine, sample had made five slides containing five serial sections each, which were stained with hematoxylin-eosin (H & E) and analysed under a light microscope. The material was analysed at the Veterinary Pathology Laboratory of the Federal University of Mato Grosso do Sul, under a Zeiss optical microscope, coupled to a microcomputer with a computer program Motic Images Plus 2.0 ML.

For intestinal histomorphometric analyses, 25 intestinal villi per animal were measured, totaling 300 villi per treatment (Schamber 2008; Silva et al. 2010) in a computer program Motic Images Plus 2.0 ML.

Bacterial challenge

The bacterial challenge was performed at the end of the probiotic supplementation period. The fish were anesthetized in eugenol, and then, the bacterium *A. hydrophila* was inoculated intraperitoneally at a concentration of 2×10^8 CFU mL^{-1} according to Silva et al. (2012). A total of 90 animals were challenged, with 18 fish per treatment. Mortality and clinical signs of bacteriosis were observed and recorded every 24 h.

The *Aeromonas hydrophila* strain (ATCC 7966) used in the bacterial challenge was supplied by the Marine Shrimps Laboratory, Department of Aquaculture, Federal University of Santa Catarina. This strain was obtained during a bacterial outbreak with high mortality in a fish farm and was isolated from symptomatic surubim, presenting haemorrhagic septicemia. To prepare the material used in the experimental challenge, *A. hydrophila* bacteria were sown in 50 mL BHI (brain heart infusion) and incubated for 24 h in a bacteriological incubator at 30 °C. Then, it was centrifuged at 400g for 20 min, the supernatant was discarded and the pellet resuspended in sterile saline (0.65% sodium chloride) to maintain bacterial concentration. For each gram of fish, 0.01 mL of this bacterial suspension was used. Moribund and dead fish were removed from the tanks daily. To calculate the relative percent survival (RPS), the formula $RPS = (1 - (\% \text{ mortality} / \% \text{ mortality control})) \times 100$ was used, according to Amend (1981).

Statistical analysis

The experiment was structured presenting a logical ordering between the treatments that can be expressed as a function of each other. Regression analysis was used to verify the variation of y as a function of x . Data were submitted to analysis of variance (ANOVA) using Statistica 13.0 (Statsoft Inc., Tulsa, USA). Tukey's test was used to compare means, with significance level of 5%. Data transformations were used according to pertinence.

Results

Growth performance

During the experimental period, the fish readily accepted diets containing probiotic *B. subtilis* and showed no behavioral changes. Weight gain (WG) was statistically different ($p < .05$) in fish in groups 1%, 2%, 3% and 4% when compared with fish in control group. Feed intake (FI) was lower in fish from the control group ($p < .05$) when compared with fish from other treatments. Among the four levels of inclusion of the probiotic, feed intake remained the same. Likewise, the biomass gain (BG) was statistically different ($p < .05$) between the groups. The fish in the 3% and 4% group obtained greater biomass gain when compared with the fish in the control group and similar to the other supplemented groups. In the other productive indexes, there were no statistical differences between the treatments (Table 2).

Haematological parameters

A statistical difference ($p < .05$) was observed in the haematological parameters of the fish between the treatments within the sampling periods, as well as between the sampling periods.

The most expressive results were observed on the 20th day of supplementation, where a number of monocytes in the fish of the 2%, 3% and 4% groups were statistically different when compared with the control group fish. Eosinophils were statistically higher in fish in the 3% group when compared with the other groups. Leukocyte numbers were statistically higher in fish in group 1% when compared with fish in control group and lymphocyte number was higher in fish in group 4% when compared with fish in group 2% and 3%, but similar to fish in

Table 2 Productive indexes. Mean values (\pm standard deviation) of final weight (FW); weight gain (WG); apparent feed conversion (AFC); feed intake (FI); biomass gain (BG); protein efficiency rate (PER) and specific growth rate (SGR) of *Pseudoplatystoma* sp. fed probiotic containing *B. subtilis* for 20 days. Different letters in the column indicate significant difference between groups ($p < .05$)

Productive indexes	Control	Inclusion levels			
		1%	2%	3%	4%
FW (g)	26.88 \pm 1.29	28.31 \pm 2.04	27.75 \pm 1.21	28.98 \pm 0.95	28.50 \pm 2.14
WG (g)	2.03 \pm 1.11b	3.85 \pm 1.01a	3.90 \pm 0.67a	4.96 \pm 0.98 ^a	4.66 \pm 0.76a
AFC	4.48 \pm 2.43	3.07 \pm 1.21	2.98 \pm 0.93	2.56 \pm 1.17	2.84 \pm 1.69
FI (g)	8.05 \pm 0.15b	9.68 \pm 0.16 ^a	10.00 \pm 0.15 ^a	10.07 \pm 0.81 ^a	9.85 \pm 0.24a
BG (g)	23.26 \pm 10.68b	34.65 \pm 9.02ab	35.05 \pm 6.08ab	44.59 \pm 8.79a	41.94 \pm 6.88a
PER (G)	5.82 \pm 3.03	7.39 \pm 2.52	7.28 \pm 1.88	8.99 \pm 3.12	8.55 \pm 3.01
SGR (%)	0.45 \pm 0.23	0.66 \pm 0.20	0.69 \pm 0.17	0.84 \pm 0.28	0.78 \pm 0.25

control and 1% groups. Glucose was statistically different in fish in the 4% group when compared with the other groups. It was also possible to observe a significant difference ($p < 0.05$) in the number of erythrocytes, hemoglobin and MCHC between the supplemented and non-supplemented fish.

The probiotic caused significant changes ($p < .05$) in the haemogram between the sampling periods on days 0, 10 and 20. The comparisons over time showed an activation of specialized cells of the fish's nonspecific defense mechanism. The number of erythrocytes, neutrophils, eosinophils, basophils, lymphocytes, monocytes, special granulocytic cells and leukocytes was statistically different in the collection on the 10th when compared with the collection in the basal period; however, it was the same when compared with the collection on the 20th (Table 3).

Phagocytic capacity

The inclusion of the probiotic *B. subtilis* in the diet of *Pseudoplatystoma* sp. increased phagocytic capacity ($p < .05$) in fish in the 4% group when compared with fish in the control and 1% group. However, it was similar to fish in groups 2% and 3%. There was no significant difference ($p > 0.05$) between the treatments for the phagocytic index (Fig. 1).

Intestinal histomorphometry

In the analysis of intestinal histomorphometry, the results showed significant differences ($p < .05$) between the groups. Villi height (μm) was higher in fish in group 2% when compared with fish in groups 1%, 3% and 4%, but it was similar to fish in the control group. In the measures of Villi width (μm), the fish in the 2% group was statistically greater when compared with the fish in the control group and similar to the other supplemented groups. Villi thickness (μm) was statistically higher in fish in the 1% group when compared with the other treatments (Fig. 2 and Table 4).

Bacterial challenge

Significant differences ($p < .05$) were observed in survival rates between the groups in the first 24 h of exposure to the pathogen. The fish in the 3% group had higher survival rates when compared with the fish in the control groups, 1% and 4%; however, it was similar to the fish in the 2% group. In subsequent periods in 48, 72, 96 and 120 h of exposure to *A. hydrophila*, mortality was observed in all groups, but without significant differences. After 120 h, no mortality was observed.

In absolute numbers, after 120 h of exposure to the pathogen, the fish in group 4% had the lowest percentages of survival (61.1%), followed by fish in groups 1% and 2% (72.2%). Fish not supplemented had a survival rate of 77.8%, while fish in the 3% probiotic kg of feed⁻¹ group had a survival rate of 83.3 (Fig. 3).

Discussion

The present study verified the influence of dietary supplementation containing *B. subtilis* C-3102 for juveniles *Pseudoplatystoma* sp. The results obtained in the present study offer a

Table 3 Complete haemogram of juvenile *Pseudoplatystoma* sp. after dietary supplementation with *B. subtilis*. The table shows the haematological parameters in the different sampling periods, basal (0 day), 10 and 20 days of probiotic supplementation. Lower case letters (a, b, c) in the horizontal observation represent statistical differences ($p < .05$) between treatments within the sample periods. Capital letters (A, B) in vertical observation represent statistical differences ($p < .05$) of the parameters within the groups between the sample periods

Haematological parameter	Basal			Inclusion			Levels		
	Control	1%	2%	1%	2%	3%	3%	3%	4%
Erythrocytes ($\times 10^6 \mu\text{L}^{-1}$)	1.60 ± 0.10bB	1.59 ± 0.12bB	1.67 ± 0.16b	1.59 ± 0.12bB	1.67 ± 0.16b	2.38 ± 0.60a	2.38 ± 0.60a	1.79 ± 0.10b	4%
Neutrophils ($\times 10^3 \mu\text{L}^{-1}$)	8.68 ± 1.74bB	11.31 ± 3.10abB	7.41 ± 1.05bB	11.31 ± 3.10abB	7.41 ± 1.05bB	18.38 ± 2.86aB	18.38 ± 2.86aB	14.88 ± 1.81abB	14.88 ± 1.81abB
Eosinophils ($\times 10^3 \mu\text{L}^{-1}$)	0.09 ± 0.03b	0.06 ± 0.03b	0.09 ± 0.03b	0.06 ± 0.03b	0.09 ± 0.03b	0.32 ± 0.11aB	0.32 ± 0.11aB	0.22 ± 0.09abAB	0.22 ± 0.09abAB
Basophils ($\times 10^3 \mu\text{L}^{-1}$)	0.06 ± 0.03bB	0.11 ± 0.07ab	0.08 ± 0.01b	0.11 ± 0.07ab	0.08 ± 0.01b	0.24 ± 0.06ab	0.24 ± 0.06ab	0.28 ± 0.05aAB	0.28 ± 0.05aAB
Lymphocytes ($\times 10^3 \mu\text{L}^{-1}$)	12.68 ± 1.96cB	10.36 ± 2.7cB	5.98 ± 1.11cB	10.36 ± 2.7cB	5.98 ± 1.11cB	22.78 ± 3.73abB	22.78 ± 3.73abB	23.77 ± 2.88aB	23.77 ± 2.88aB
Monocytes ($\times 10^3 \mu\text{L}^{-1}$)	0.08 ± 0.02	0.13 ± 0.07	0.06 ± 0.02	0.13 ± 0.07	0.06 ± 0.02	0.04 ± 0.02B	0.04 ± 0.02B	0.09 ± 0.04B	0.09 ± 0.04B
Special granulocytic cells ($\times 10^3 \mu\text{L}^{-1}$)	0.21 ± 0.04ab	0.41 ± 0.11ab	0.10 ± 0.03bB	0.41 ± 0.11ab	0.10 ± 0.03bB	0.65 ± 0.28a	0.65 ± 0.28a	0.49 ± 0.13ab	0.49 ± 0.13ab
Leukocytes ($\times 10^3 \mu\text{L}^{-1}$)	22.16 ± 3.67bB	22.75 ± 6.11bB	13.91 ± 2.19bB	22.75 ± 6.11bB	13.91 ± 2.19bB	42.94 ± 6.53aB	42.94 ± 6.53aB	40.39 ± 4.78aB	40.39 ± 4.78aB
Thrombocytes ($\times 10^5 \mu\text{L}^{-1}$)	35.55 ± 12.36	27.57 ± 5.97A	19.60 ± 2.76A	27.57 ± 5.97A	19.60 ± 2.76A	24.45 ± 3.73A	24.45 ± 3.73A	23.56 ± 3.21A	23.56 ± 3.21A
Haemoglobin (g dL ⁻¹)	na	na	na	na	na	4.5 ± 0.28B	4.5 ± 0.28B	4.5 ± 0.35AB	4.5 ± 0.35AB
Haematocrit (%)	26.75 ± 2.25	27.00 ± 1.87	26.67 ± 1.38	27.00 ± 1.87	26.67 ± 1.38	25.41 ± 1.24	25.41 ± 1.24	26.08 ± 1.38AB	26.08 ± 1.38AB
MCV (fL)	170.04 ± 13.01A	175.38 ± 10.65A	146.93 ± 10.78	175.38 ± 10.65A	146.93 ± 10.78	190.32 ± 37.80	190.32 ± 37.80	149.05 ± 10.22	149.05 ± 10.22
MCHC (g dL ⁻¹)	na	na	na	na	na	15.29 ± 2.33AB	15.29 ± 2.33AB	17.75 ± 0.94A	17.75 ± 0.94A
Glucose (mg dL ⁻¹)	42.96 ± 8.70abA	46.27 ± 2.34aa	41.75 ± 3.31abA	46.27 ± 2.34aa	41.75 ± 3.31abA	34.27 ± 3.94ab	34.27 ± 3.94ab	29.96 ± 5.39bB	29.96 ± 5.39bB
10 days	Control	1%	2%	1%	2%	3%	3%	4%	4%
Erythrocytes ($\times 10^6 \mu\text{L}^{-1}$)	2.25 ± 0.10aA	2.10 ± 0.18abA	1.64 ± 0.06c	2.10 ± 0.18abA	1.64 ± 0.06c	1.93 ± 0.06abc	1.93 ± 0.06abc	1.65 ± 0.09bc	1.65 ± 0.09bc
Neutrophils ($\times 10^3 \mu\text{L}^{-1}$)	40.50 ± 5.70A	51.11 ± 10.10A	36.76 ± 3.45A	51.11 ± 10.10A	36.76 ± 3.45A	46.29 ± 5.19A	46.29 ± 5.19A	37.54 ± 4.69A	37.54 ± 4.69A
Eosinophils ($\times 10^3 \mu\text{L}^{-1}$)	0.34 ± 0.11	0.28 ± 0.13	0.31 ± 0.16	0.28 ± 0.13	0.31 ± 0.16	0.18 ± 0.08B	0.18 ± 0.08B	0.06 ± 0.06B	0.06 ± 0.06B
Basophils ($\times 10^3 \mu\text{L}^{-1}$)	0.45 ± 0.13A	0.44 ± 0.12	0.16 ± 0.07	0.44 ± 0.12	0.16 ± 0.07	0.44 ± 0.12	0.44 ± 0.12	0.63 ± 0.30A	0.63 ± 0.30A
Lymphocytes ($\times 10^3 \mu\text{L}^{-1}$)	37.47 ± 3.61A	53.92 ± 6.30A	41.35 ± 28.31A	53.92 ± 6.30A	41.35 ± 28.31A	54.59 ± 7.98A	54.59 ± 7.98A	42.77 ± 5.27AB	42.77 ± 5.27AB
Monocytes ($\times 10^3 \mu\text{L}^{-1}$)	0b	0.03 ± 0.03b	0.11 ± 0.06ab	0.03 ± 0.03b	0.11 ± 0.06ab	0.28 ± 0.14aB	0.28 ± 0.14aB	0bB	0bB
Special granulocytic cells ($\times 10^3 \mu\text{L}^{-1}$)	0.22 ± 0.10	0.58 ± 0.20	0.47 ± 0.15A	0.58 ± 0.20	0.47 ± 0.15A	0.61 ± 0.19	0.61 ± 0.19	0.43 ± 0.17	0.43 ± 0.17
Leukocytes ($\times 10^3 \mu\text{L}^{-1}$)	82.73 ± 6.55A	107.51 ± 15.46A	80.04 ± 5.08A	107.51 ± 15.46A	80.04 ± 5.08A	103.82 ± 7.92A	103.82 ± 7.92A	82.73 ± 6.55A	82.73 ± 6.55A
Thrombocytes ($\times 10^5 \mu\text{L}^{-1}$)	12.93 ± 4.41	10.20 ± 5.75B	3.69 ± 1.40B	10.20 ± 5.75B	3.69 ± 1.40B	9.82 ± 4.30AB	9.82 ± 4.30AB	5.43 ± 2.57B	5.43 ± 2.57B
Haemoglobin (g dL ⁻¹)	5.03 ± 0.21	4.55 ± 0.31A	4.84 ± 0.31	4.55 ± 0.31A	4.84 ± 0.31	5.03 ± 0.28A	5.03 ± 0.28A	4.77 ± 0.22A	4.77 ± 0.22A
Haematocrit (%)	29.17 ± 0.71a	27.08 ± 1.33ab	24.33 ± 1.19b	27.08 ± 1.33ab	24.33 ± 1.19b	25.33 ± 1.14ab	25.33 ± 1.14ab	23.67 ± 1.01bB	23.67 ± 1.01bB
MCV (fL)	133.00 ± 7.49B	134.74 ± 7.38B	149.61 ± 7.14	134.74 ± 7.38B	149.61 ± 7.14	132.33 ± 6.14	132.33 ± 6.14	145.67 ± 8.63	145.67 ± 8.63
MCHC (g dL ⁻¹)	17.26 ± 0.68ab	16.78 ± 0.88bA	20.43 ± 1.66a	16.78 ± 0.88bA	20.43 ± 1.66a	19.94 ± 0.89abA	19.94 ± 0.89abA	20.34 ± 0.94aB	20.34 ± 0.94aB
Glucose (mg dL ⁻¹)	15.97 ± 3.02bB	20.91 ± 5.47bB	19.76 ± 4.31bB	20.91 ± 5.47bB	19.76 ± 4.31bB	28.05 ± 4.64ab	28.05 ± 4.64ab	35.79 ± 4.38aB	35.79 ± 4.38aB

Table 3 (continued)

	Basal				Inclusion				Levels				
	Control	1%	2%	3%	Control	1%	2%	3%	Control	1%	2%	3%	4%
20 days													
Erythrocytes ($\times 10^6 \mu\text{L}^{-1}$)	2.07 \pm 0.13aA	2.06 \pm 0.07aA	1.70 \pm 0.11b	1.84 \pm 0.11ab	2.07 \pm 0.13aA	2.06 \pm 0.07aA	1.70 \pm 0.11b	1.84 \pm 0.11ab	2.07 \pm 0.13aA	2.06 \pm 0.07aA	1.70 \pm 0.11b	1.84 \pm 0.11ab	1.87 \pm 0.09ab
Neutrophils ($\times 10^3 \mu\text{L}^{-1}$)	32.58 \pm 3.94abA	38.39 \pm 2.23aA	25.32 \pm 1.52bcA	18.14 \pm 2.37cB	32.58 \pm 3.94abA	38.39 \pm 2.23aA	25.32 \pm 1.52bcA	18.14 \pm 2.37cB	32.58 \pm 3.94abA	38.39 \pm 2.23aA	25.32 \pm 1.52bcA	18.14 \pm 2.37cB	22.1 \pm 0.2.13cB
Eosinophils ($\times 10^3 \mu\text{L}^{-1}$)	0.15 \pm 0.11b	0b	0.13 \pm 0.13b	1.46 \pm 0.47aA	0.15 \pm 0.11b	0b	0.13 \pm 0.13b	1.46 \pm 0.47aA	0.15 \pm 0.11b	0b	0.13 \pm 0.13b	1.46 \pm 0.47aA	0.60 \pm 0.18bA
Basophils ($\times 10^3 \mu\text{L}^{-1}$)	0.15 \pm 0.11AB	0.12 \pm 0.08	0.08 \pm 0.08	0.26 \pm 0.11	0.15 \pm 0.11AB	0.12 \pm 0.08	0.08 \pm 0.08	0.26 \pm 0.11	0.15 \pm 0.11AB	0.12 \pm 0.08	0.08 \pm 0.08	0.26 \pm 0.11	0b
Lymphocytes ($\times 10^3 \mu\text{L}^{-1}$)	45.76 \pm 5.22abA	47.19 \pm 7.21abA	29.05 \pm 4.08bA	43.78 \pm 5.16abAB	45.76 \pm 5.22abA	47.19 \pm 7.21abA	29.05 \pm 4.08bA	43.78 \pm 5.16abAB	45.76 \pm 5.22abA	47.19 \pm 7.21abA	29.05 \pm 4.08bA	43.78 \pm 5.16abAB	54.67 \pm 5.22aA
Monocytes ($\times 10^3 \mu\text{L}^{-1}$)	0.12 \pm 0.08ab	0 \pm 0b	0.87 \pm 0.61ab	1.03 \pm 0.21aA	0.12 \pm 0.08ab	0 \pm 0b	0.87 \pm 0.61ab	1.03 \pm 0.21aA	0.12 \pm 0.08ab	0 \pm 0b	0.87 \pm 0.61ab	1.03 \pm 0.21aA	0.97 \pm 0.25aA
Special granulocytic cells ($\times 10^3 \mu\text{L}^{-1}$)	0.33 \pm 0.33	0.58 \pm 0.52	0.28 \pm 0.24AB	0.82 \pm 0.22	0.33 \pm 0.33	0.58 \pm 0.52	0.28 \pm 0.24AB	0.82 \pm 0.22	0.33 \pm 0.33	0.58 \pm 0.52	0.28 \pm 0.24AB	0.82 \pm 0.22	0.83 \pm 0.37
Leukocytes ($\times 10^3 \mu\text{L}^{-1}$)	79.67 \pm 8.55abA	86.75 \pm 7.93aA	56.80 \pm 5.13bA	66.01 \pm 6.03abB	79.67 \pm 8.55abA	86.75 \pm 7.93aA	56.80 \pm 5.13bA	66.01 \pm 6.03abB	79.67 \pm 8.55abA	86.75 \pm 7.93aA	56.80 \pm 5.13bA	66.01 \pm 6.03abB	80.01 \pm 12.74abA
Thrombocytes ($\times 10^3 \mu\text{L}^{-1}$)	11.95 \pm 5.77	10.55 \pm 3.70AB	9.27 \pm 3.05AB	10.31 \pm 3.36B	11.95 \pm 5.77	10.55 \pm 3.70AB	9.27 \pm 3.05AB	10.31 \pm 3.36B	11.95 \pm 5.77	10.55 \pm 3.70AB	9.27 \pm 3.05AB	10.31 \pm 3.36B	5.45 \pm 2.53B
Haemoglobin (g dL ⁻¹)	6.39 \pm 0.70a	3.74 \pm 0.19bB	4.49 \pm 0.49b	3.89 \pm 0.27bB	6.39 \pm 0.70a	3.74 \pm 0.19bB	4.49 \pm 0.49b	3.89 \pm 0.27bB	6.39 \pm 0.70a	3.74 \pm 0.19bB	4.49 \pm 0.49b	3.89 \pm 0.27bB	3.91 \pm 0.25bB
Haematocrit (%)	28.67 \pm 0.76	27.58 \pm 0.80	26.58 \pm 1.37	26.67 \pm 1.03	28.67 \pm 0.76	27.58 \pm 0.80	26.58 \pm 1.37	26.67 \pm 1.03	28.67 \pm 0.76	27.58 \pm 0.80	26.58 \pm 1.37	26.67 \pm 1.03	28.42 \pm 1.07A
MCV (fL)	143.49 \pm 7.69AB	134.95 \pm 4.24B	162.34 \pm 7.91	148.46 \pm 7.15	143.49 \pm 7.69AB	134.95 \pm 4.24B	162.34 \pm 7.91	148.46 \pm 7.15	143.49 \pm 7.69AB	134.95 \pm 4.24B	162.34 \pm 7.91	148.46 \pm 7.15	153.91 \pm 7.69
MCHC (g dL ⁻¹)	22.85 \pm 2.95a	13.54 \pm 0.40bB	17.41 \pm 2.29ab	14.58 \pm 0.85bB	22.85 \pm 2.95a	13.54 \pm 0.40bB	17.41 \pm 2.29ab	14.58 \pm 0.85bB	22.85 \pm 2.95a	13.54 \pm 0.40bB	17.41 \pm 2.29ab	14.58 \pm 0.85bB	13.86 \pm 0.88bB
Glucose (mg dL ⁻¹)	41.91 \pm 6.79bA	39.41 \pm 9.78bAB	48.55 \pm 8.42bA	48.14 \pm 11.37b	41.91 \pm 6.79bA	39.41 \pm 9.78bAB	48.55 \pm 8.42bA	48.14 \pm 11.37b	41.91 \pm 6.79bA	39.41 \pm 9.78bAB	48.55 \pm 8.42bA	48.14 \pm 11.37b	95.47 \pm 19.09aA

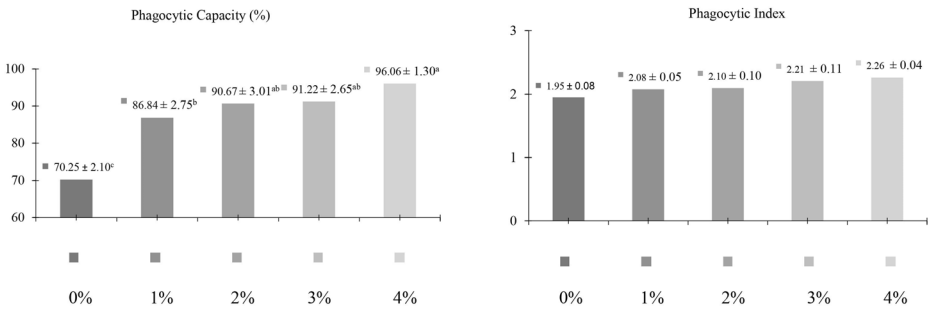


Fig. 1 Phagocytic capacity and phagocytic index of juvenile *Pseudoplatystoma* sp. supplemented for 20 days with probiotic *B. subtilis* at 1%, 2%, 3% and 4% kg feed⁻¹ and control group. Different letters indicate significant differences between the groups ($p < 0.05$)

starting point for new productive perspectives, including the haemato-immunological and sanitary aspects, as well as, important information about the influence of the probiotic on the intestinal morphology of *Pseudoplatystoma* sp.

Probiotic bacteria have been shown to contribute to fish growth. The literature proves that *Bacillus* species have probiotic properties and, when administered to fish, increase growth and improve the feed intake. Generally, the fact is related to the improvements conferred by the probiotic in the intestinal morphology, increasing the activity of digestive enzymes and,

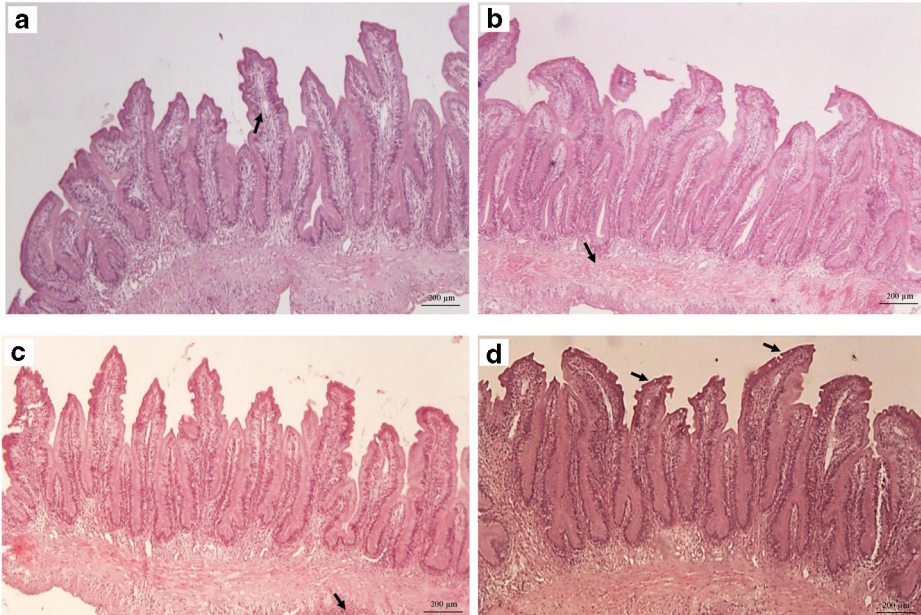


Fig. 2 Intestinal histology analysis of juvenile *Pseudoplatystoma* sp. after 20 days of dietary supplementation with probiotic *B. subtilis*. **A** The intestinal morphology of the fish in the 2% group where the highest villi height was observed (long villi and lamina propria—arrow). **B** Morphometry of the fish in the control group that presented similar villi height to the fish in the 2% group (inner muscular layer—arrow). **C** It shows the intestinal morphology of the fish from the 2% group that presented greater villi width (external transverse muscular layer—arrow). Finally **D** we can observe the intestinal morphology of the fish in the 1% group that presented greater villi thickness (brush border—arrow). HE (bar = 200 μm)

Table 4 Intestinal histomorphometry. Measurements of the midgut portion of surubins *Pseudoplatystoma* sp. after 20 days of feeding with probiotic *Bacillus subtilis* (mean ± standard error). Different letters indicate significant differences between the groups ($p < .05$)

Intestine histomorphometry	Inclusion Levels				
	Control	1%	2%	3%	4%
Villi height (µm)	271.25 ± 3.82ab	265.75 ± 3.67b	278.22 ± 2.69a	254.10 ± 2.30c	262.43 ± 2.05bc
Villi width (µm)	65.36 ± 4.26b	79.13 ± 3.55ab	85.33 ± 6.21a	73.96 ± 2.66ab	73.90 ± 2.19ab
Villi thickness (µm)	31.37 ± 1.07c	40.59 ± 0.75a	37.19 ± 0.62b	33.26 ± 0.49c	31.59 ± 0.64c

consequently, they can increase the digestive capacity of the feeds offered to fish. This is one of the reasons for choosing *Bacillus* as probiotic, because they are capable of this modulation in the digestive tract of fish (Dawood et al. 2019; Kuebutomye et al. 2019; Olmos et al. 2020).

The diets containing the probiotic showed different responses in growth performance and feed intake parameters of the present research. Likewise, Veiga et al. (2020), when offering a diet containing *B. subtilis* to *Pseudoplatystoma* sp. for 10 days, observed increases in biomass gain in fish supplemented with 3% probiotic per kilo of feed. Similarly, in the present research, fish that received the inclusion of 3% probiotic in the diet showed improvements in production performance.

According to Olmos et al. (2020) *B. subtilis* produce enzymes capable of improving the assimilation of nutrients, favoring the metabolism of carbohydrates, lipids and proteins in aquatic animals. Probiotic bacteria can facilitate the digestion of dietary proteins, as enzymes produced by probiotic bacteria can complement protease activity by increasing food digestibility (Ochoa-Solano and Olmos-Soto 2006; Kuebutomye et al. 2019). In this context, the catfish *Pseudoplatystoma* sp. as a carnivorous species (Honorato et al. 2015) could obtain productive improvements by including *B. subtilis* in its diet, since the digestion of alternative protein sources, such as those of vegetable origin, could be optimized. Replacement of even a small fraction of dietary protein may represent significant financial effectiveness and significantly lower environmental impact on large-scale production (Martino et al. 2002).

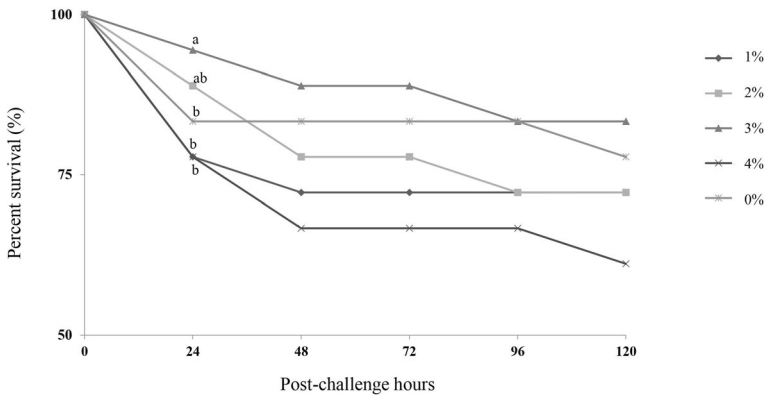


Fig. 3 Percent survival of juvenile *Pseudoplatystoma* sp. supplemented for 20 days with probiotic *B. subtilis* at 1%, 2%, 3% and 4% kg feed⁻¹ and control (0%) group after challenge with *Aeromonas hydrophila*. Different letters indicate significant differences ($p < .05$) in survival rates between the groups at the respective analysis times

Haematological analyses may reflect the sanitary state of the fish. Complete fish blood cell counting is an important and powerful diagnostic tool, as well as a component of a valuable database. It can be used to monitor the health status of fish in response to changes related to nutrition, water quality and disease and in response to therapeutic methods (Fazio 2019). In the present study, three specific moments were decisive in the blood assessment of fish and, in this way, different haemograms were made. It is noted that in the present study, even without the influence of the treatment applied to the fish, the blood parameters may show marked variations, as observed in the basal data. If we observe the haemogram of the basal samples, there were differences in the number of erythrocytes between the groups. This fact may indicate that some stressful factor affected the fish prior to collection and reflecting in the other haematological indexes of the basal collection.

Stress is one of the factors that can make fish susceptible to disease, causing changes in the innate immune responses that occur at the cellular level in fish. In aquaculture, stressors are part of the activity characteristics, where handling, water quality, transport and stocking densities can affect fish physiology and health (Kuebutornye et al. 2019). The use of a probiotic based on *B. subtilis* was able to provide a stress reduction in *Clarias gariepinus* catfish, making it an important tool to mitigate the negative effects of cultivation systems (Romanova et al. 2020).

The haematological results found in the present study are similar to those found by Mouriño et al. (2016), when they performed a hematological approach of the hybrid surubim *Pseudoplatystoma reticulatum* × *P. corruscans*. At the time of the research, they investigated bacterial strains with probiotic potential isolated from fish intestines. Compared with control fish, fish supplemented with probiotic *W. cibaria* showed an increase in RBC. In contrast, Veiga et al. (2020), when supplementing the diets of *Pseudoplatystoma* sp. with *B. subtilis* for 10 days, did not observe significant differences for haematocrit, hemoglobin, erythrocytes, MVC, MCHC, glucose, monocytes, basophils, eosinophils and special granulocytic cell.

Unlike in the present study, haematological evaluation on the tenth day revealed a higher number of leukocytes circulating in the blood of fish. The probable answer to this contrast observed between the two studies may be the difference in the average water temperature in the different studies, where Veiga et al. (2020) observed that the water temperature remained at an average of 26.2 ± 2.9 °C, while in the present study, it was close to 24.8 ± 1.8 °C. The thermal comfort temperature for surubim is close to 28 °C (Campos 2005). Thus, it is very likely that, in the present study, the probiotic triggered a more consistent non-specific response on the tenth day of the experiment due to possible thermal stress due to the lower temperature.

The results of the haemogram of the present study showed that the probiotic supplementation was able to activate some leukocyte lineages in a representative manner between the groups. The cells that support the fish's non-specific immune system are circulating in the bloodstream and immune responses can be triggered at any time in response to physical injuries or inflammatory processes. In addition, the phagocytic activities of B lymphocytes, granular eosinophilic cells present in the intestinal mucosa and gills can respond to bacteria and parasites (Ranzani-Paiva and Silva-Souza 2004; Kiron 2012).

The phagocytic process and cellular metabolism end up resulting in the production of reactive oxygen species that can be harmful when the production of lipid peroxidation exceeds the antioxidant capacity of cells or tissues. In these cases, *Bacillus* spp., such as probiotics, can produce antioxidant enzymes, for example superoxide dismutase and glutathione, to effectively eliminate free radicals. Whether in serum or mucus, *Bacillus* spp. can modulate antioxidant activities (Kuebutornye et al. 2019). The inclusion of *B. subtilis* in juvenile *Pseudoplatystoma*

sp. diets provided higher phagocytic activity in all treated groups. Immune responses involving phagocytosis are assisted by non-specific immune cells such as monocytes/macrophages, neutrophils, among others. Phagocytosis is one of the main mechanisms used to remove pathogens and cellular debris (Kiron 2012). Thus, we can infer that the probiotic treatment had a positive effect on the fish immune system, increasing the innate defense capacity of the animals that received the additive at different inclusion levels.

The gut is a multifunctional organ directly involved in nutrition, so it is strongly related to fish immunity (Sugita et al. 1998; Kiron 2012). Cell renewal through mitosis or loss of cells at the base of the villi is caused by the development of the intestinal mucosa, increasing the density and height of the epithelial cells (Maiorka et al. 2002). In these cases of cell renewal, through increased cell proliferation, nutrient digestion and absorption are maximized with greater weight gain (Boleli et al. 2000).

In the present study, we observed a reduction in villi height at the highest levels of inclusion of *B. subtilis* (3% and 4%), suggesting that the gradual increase in the probiotic levels did not cause the development of the intestinal mucosa. These results corroborate with Veiga et al. (2020), who in a short communication reported that the probiotic *B. subtilis* improved the intestinal absorption surface of the hybrid *Pseudoplatystoma corruscans* × *P. reticulatum*. On the occasion, the researchers found the highest villi height in the inclusion content at 1% and the lowest heights in the highest probiotic inclusion levels.

The beneficial effects of probiotic *B. subtilis* on intestinal morphometry are related to the reduction of colonization of the intestinal epithelium by bacteria harmful to the mucosa, which act by inhibiting and competing for the adhesion sites in the enterocytes through the connection with the glycocalyx (Zhao et al. 2012; He et al. 2013; Zhou et al. 2019; Kuebutomye et al. 2020), promoting the competitive exclusion of unwanted bacteria in different segments of the intestine (Kuebutomye et al. 2019). Thus, when experimental conditions involve challenges to the immune system, it is very important to observe whether improvements in the intestinal morphology of animals will have a corresponding effect during exposure to the pathogen, as these responses are potentially related to the digestive physiology of fish when using supplements, such as probiotics.

The probiotic *B. subtilis* positively influenced the fish survival to *A. hydrophila* during the experimental challenge in the present study. Bacteria are natural components of the environment and, as such, are present in aquaculture environments. Probiotics have emerged as a sustainable solution to contain pathogens in aquaculture and bacteria of the *Bacillus* genus have been proven to more successfully combat a wide variety of fish pathogens, such as *Aeromonas* spp., *Vibrio* spp., *Streptococcus* spp., compared with other probiotics in aquaculture (Kavitha et al. 2018; Kuebutomye et al. 2020; Hayatgheib et al. 2020). In this way, *B. subtilis* is strengthened as a sustainable alternative to the use of antibiotics in aquaculture.

Tang et al. (2019) found that the dietary supplementation of probiotic *B. subtilis* affects antioxidant defenses and the immune response in grass carp exposed to *A. hydrophila*. The authors consider that *B. subtilis* can provide effective protection to fish against damage caused by *A. hydrophila* infection. Similarly, Zhang et al. (2019) found that *B. subtilis* increased intestinal apoptosis of grass carp *Ctenopharyngodon idella* challenged orally with *A. hydrophila*, suggesting that *B. subtilis* may play an important role in reducing intestinal damage after challenge.

As noted in the present study, as well as in the published literature, *Bacillus* spp. is widespread and tested on a wide variety of aquaculture species. Therefore, it becomes a viable alternative for optimizing food absorption and increased zootechnical performance, better

responses to stress conditions, increased immune response and disease resistance (Kuebutornye et al. 2019).

In conclusion, the dietary supplementation with probiotic *Bacillus subtilis* applied in the present study in the inclusions of 1%, 2%, 3% and 4% caused significant improvements in the physiology of juvenile *Pseudoplatystoma* sp. and based on the data obtained in the research, we indicate the 2% and 3% inclusions as more efficient due to the improvements observed in the productive indexes, in the intestinal morphometry, as well as, in the non-specific immunity, optimizing the survival of the animals exposed to the pathogen.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

Ethical approval All applicable international, national and/or institutional guidelines for the care and use of animals were followed.

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