



First report of *Streptococcus agalactiae* isolated from *Oreochromis niloticus* in Piura, Peru: Molecular identification and histopathological lesions

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ABSTRACT

The aim of this study was to identify the bacterium *Streptococcus agalactiae* isolated in farmed Nile tilapia (*Oreochromis niloticus*) from Piura, Peru and to characterize the histopathological lesions caused by this pathogen. Sixteen tilapias were sampled with clinic signs of the disease such as erratic swimming, exophthalmia and haemorrhages on the body and fins. Qualitative PCR in real time and histopathological analysis were performed. Nine fishes positives to *S. agalactiae* were found. The main histopathological findings were fibrinosuppurative epicarditis, periesplenitis, meningitis and panophthalmitis with predominance of mononuclear infiltration in all tissues. The correlation between qPCR and histopathological findings demonstrated nine fish (prevalence of 56.25%) with Cq lower than 30, associated to high degree of tissue injuries. This study reports the first isolation of *S. agalactiae* by PCR in real time in tilapia farmed in Peru and characterizes the major histopathological changes caused by this bacterium.

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1. Introduction

The tilapia industry has positive prospects for productive development in Peru (Sandoval et al., 2010). The country has about 154,000 ha available to develop tilapia farms in earthen ponds, concrete or geomembranes (Baltazar, 2009). The departments of San Martín and Piura have suitable areas for the production of this species (PRODUCE, 2011), because the waters offer exceptional conditions for breeding, such as warm temperatures (20–30 °C), low pollution, turbidity and good oxygenation (Baltazar and Palomino, 2004).

The culture of Nile tilapia is one of the fastest growing aquaculture activities worldwide (Conroy et al., 2008). However, the intensification of fish production has resulted in the occurrence of

stress (Belo et al., 2005, 2012), a fact that increases the susceptibility to diseases outbreaks, such as bacteria (Amal and Zamri-Saad, 2011), parasites (Manrique et al., 2012, 2015a, 2016; Padua et al., 2015), nutritional deficiencies (Sakabe et al., 2013; Belo et al., 2014; Castro et al., 2014a) and fungi (Jiménez, 2010; Iregui et al., 2014). Among them, bacterial are the main obstacle to the sustainable development of the culture of tilapia (Yuasa et al., 2008).

The genus *Streptococcus* is one of the most important Gram-positive bacteria that affect the culture of tilapia (Conroy, 2009; Jiménez, 2010). Within this, the species *S. agalactiae* is the most prevalent worldwide (Sheehan et al., 2009). *S. agalactiae* is a coccus, organized in pairs or short, catalase and oxidase negative and positive CAMP chains; it may or may not be haemolytic (Abuseliana et al., 2010; Buller, 2014). The lesions are characterized by loss of appetite, exophthalmia, eye haemorrhage, corneal opacity, distended abdomen, curvature of the spinal cord, erratic swimming, stiffness and bleeding on the base of the fins (Yanong and Francis-Floyd, 2002; Pulido and Iregui, 2010). Castro et al. (2014b) studied

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the Kinect of cell accumulation in swim bladder of tilapias during acute aerocystitis by *S. agalactiae* and verified high number of granulocytes in site of inflammation, while Pulido and Iregui (2010) described infiltration of mononuclear cells with subsequent granuloma formation and meningoencephalitis in chronic inflammatory response.

The diagnosis of *S. agalactiae* is based on clinical signs, histopathological lesions and microbiological isolation (Wongsathein, 2012). However, the culture methods and biochemical tests can be confused for the variable reactions of the different strains of *S. agalactiae* (Wongsathein, 2012). For this, several authors suggest the specific identification of this bacterium through molecular techniques such as PCR (Jimenez et al., 2011).

Tilapias reared in Piura, the main production area in Peru, have presented clinical signs and lesions described in outbreaks of *S. agalactiae* (Sedano, 2013). However, the presence of this bacterium has not been confirmed so far. The objective of this study was to identify by PCR in real time the presence of *S. agalactiae* and characterize histopathological lesions in diseased tilapias from nursery and grow-out phases.

2. Materials and methods

2.1. Study design

The study design was cross-sectional. Eight tilapias in each of the phases nursery (33.5 ± 3.6 g) and grow-out (670.5 ± 22.6 g) were selected between February and June 2014. The criterion of selection of individuals was the presence of clinical signs such as erratic swimming, exophthalmia, lethargy, bleeding around the body and superficial location in the water column. The fish were euthanized with a lethal dose of sodium benzocaine 1:500 (v/v) previously diluted in pure alcohol (0.1 g/mL) (Manrique et al., 2015b), then proceeded to the necropsy according to the protocol described by Ferguson (2006).

The sampling was performed aseptically for the microbiology analysis of the following organs: spleen, brain, liver and kidney. For histopathology, besides were taken samples of gills, heart, eye and skeletal muscle of approximately 1.0 cm^2 and fixed in 10% buffered formalin.

2.2. Location

The fish farm was located in the department of Piura ($4^\circ 38'27''\text{S}$, $80^\circ 32'55''\text{W}$), Northern Peru. The water surface area was 19.26 ha, supplied by a water reserve of the valley. The tanks were concreted with stoking density of 15–19 kg of biomass by m^3 of water. The study was conducted between February and June 2014, ambient temperatures ranging from 24 to 35 °C. The average values of the water quality parameters reported by the fish farm during the period can be seen in Table 1.

2.3. Microbiology

Samples of spleen, brain, liver and kidney were plated on blood agar (enriched with 5% sheep blood) and incubated at 30 °C for 24–48 h. Then, there were identified the morphology of the colonies, Gram positive cocci, catalase and oxidase negative, as isolated presumptive of *Streptococcus* spp.

2.4. DNA extraction

The DNA of the isolated presumptive to *Streptococcus* spp. was extracted with the PureLink Genomic DNA Kit (Invitrogen, USA), following the manufacturer's instructions for Gram positive bacteria.

For purification of DNA, it was used the same commercial kit PureLink Genomic DNA Kits (Invitrogen, USA). The material was centrifuged at 13,000g for 1 min at room temperature (24 °C) followed by two washes with 40 mL of elution buffer in each wash. The concentration and quality of DNA extracted was measured with Nanodrop spectrophotometer 2000 UV–vis (Thermo Scientific, USA). Finally, the genetic material was stored at -20°C until use.

The positive control of the reaction was strain of *Streptococcus agalactiae*, donated by the Laboratory Genomic Medicine, University Surcolombiana – Colombia. The DNA extraction was a performed with commercial PureLink Genomic DNA Kits (Invitrogen, USA), following the manufacturer's recommendations. As negative control DEPC water was used (Invitrogen, USA).

2.5. Sensitivity and specificity of real-time PCR

Previously, tests for PCR were performed in real time with different dilutions to optimize the reaction, from a concentration of 1/1 (15 ng/ μL) until the concentration of 1/100,000. Fig. 1 shows the DNA extracted from three species of *Streptococcus*: *S. pyogenes*, *S. pneumoniae* and *S. iniae*, which shows the high specificity of the test to identify the species *S. agalactiae*.

2.6. Real-time PCR

PCR was performed in qualitative real time. Primers were designed using the intergenic fraction of the 16S–23S rRNA as amplification region, which were synthesized from the strain *S. agalactiae* SA 66-07, access number AF064441, reported in Genbank, using the Primer Express 3.0 Applied Biosystems (Life Technologies, USA) (Betancur et al., 2012) software. The primers used for the detection of *S. agalactiae* were Forward (5' CATTTCGGCTCTGT-TAGTTTTGAG 3') Reverse (5' GGAGCCTAGCGGATCGA 3') and probe (5' VIC- AGAGCGCTGCTTGCACGCA-TAMRA 3') (Bioneer Corporation, Korea).

The PCR reaction was performed in triplicate, briefly: 12.5 μL Taqman Universal PCR Master Mix (2x) (Invitrogen, USA), 0.5 μL probe (10 μM), 1.25 μL each primers (10 μM), 1 μL of DNA (20 $\mu\text{g}/\mu\text{L}$) and 8.5 μL of DEPC water (Invitrogen) to a total volume of 25 μL . Thermal conditions were: denaturation at 95 °C for 10 min followed by 40 cycles of 15 s at 95 °C, 2 min annealing at 50 °C and 1 min of extension at 60 °C, using the Applied Biosystem equipment 7300 (Life Technologies, USA).

2.7. Histopathology

Samples of spleen, gills, brain, heart, eye and liver were fixed in 10% buffered formalin and processed according to the standard protocol for histological studies of fixed tissues. Sections of thickness 3 μm were cut and mounted on slides and were stained with hematoxylin eosin (H&E).

3. Results

3.1. Necropsy

The main necropsy findings were observed: diffuse haemorrhagic areas around the mouth and fins (3) 18.75%, exophthalmos and eye opacity (4) 25%, hyphema (2) 12.5%, ascites (4) 25% congested brain (7) 43.75%, liver and spleen congestion (3) 18.75%, cloudy spots in liver and heart (12) 75%, hepatomegaly, splenomegaly and ascites (5) 31.25%, dilated gut (4) 25%, melanosis (2) 12.5%.

Table 1
Parameters of water quality (mean \pm SD) obtained in the tanks during the period of study (February to June, 2014).

	Temperature ($^{\circ}$ C)	Oxygen (mg/L)	CO ₂ (mg/L)	pH	Nitrites (mg/L)	Ammonium (mg/L)	Hardness (mg/L)
Nursery	25.4 \pm 1.1	4.2 \pm 0.8	3.1 \pm 0.5	7.3 \pm 0.4	0.0	0.14 \pm 0.03	92.5 \pm 3.2
Grow-out	25.2 \pm 0.9	4.0 \pm 0.6	3.0 \pm 0.4	5.8 \pm 0.8	0.0	0.34 \pm 0.02	93.2 \pm 2.8

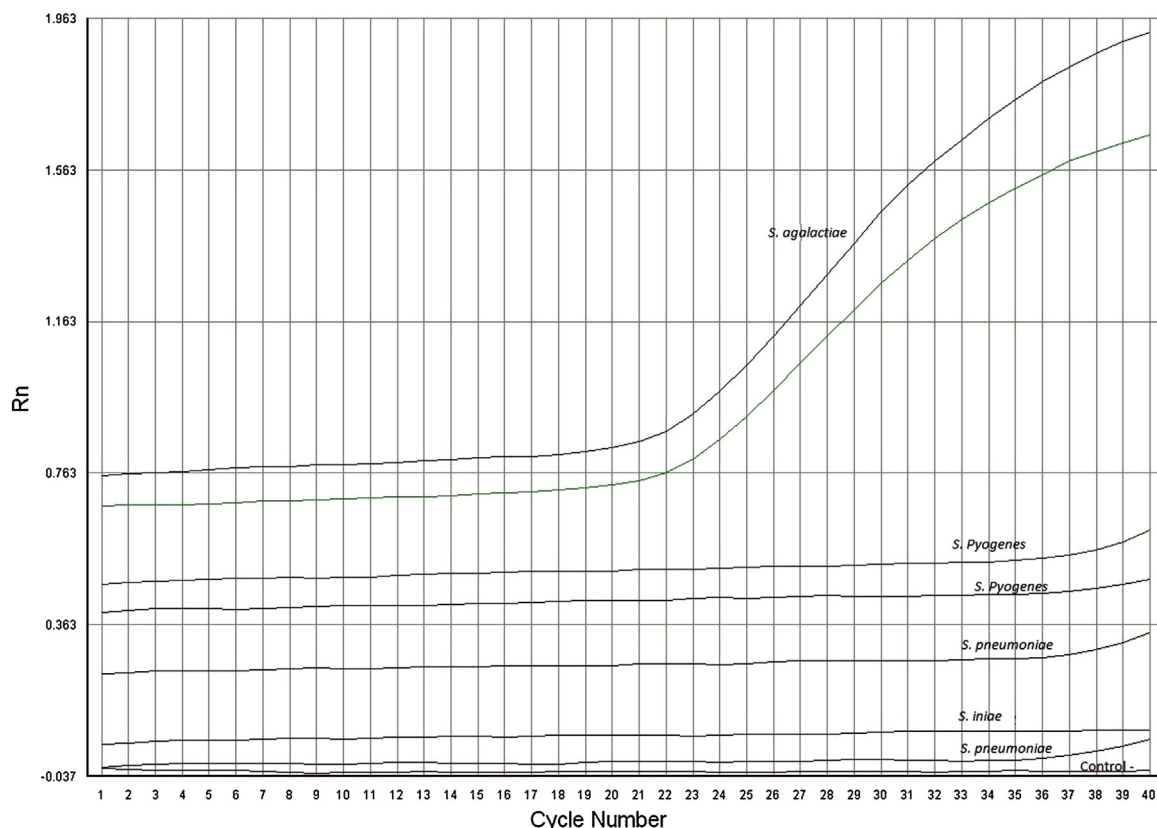


Fig. 1. Real time PCR with different species of *Streptococcus* to determinate the specificity of the assay.

3.2. Microbiology

Were identified 30 isolated presumed for *Streptococcus* spp. The distribution of the isolated by organs was the following: Liver (1) 3.33%, brain (8) 26.67%, kidney (10) 33.33% and spleen (11) 36.67%. These isolated belong to 16 tilapias positives presumed to *Streptococcus* spp.

3.3. Histopathology

The most important histopathology findings were: acute or chronic suppurative fibrin type epicarditis (11) 68.75%; suppurative myocarditis (5) 31.25%; acute suppurative meningitis (10) 62.5% and acute suppurative panophthalmitis. Presence of dispersed pigment in epithelial cells (12) 75%, coagulation necrosis in muscle (14) 87.5%, necrosis and mononuclear infiltration in the liver (9) 56.25%, hyperplasia and fusion of secondary lamella (12) 75%, congestion (5) 31.25%, increased of melanomacrophages center (MMC) in spleen (5) 31.25%, acute suppurative periesplenitis (1) 6.25%, lymphocytolysis (6) 37.50%, fat and hydropic degeneration (8) 50% (Fig. 2–5).

3.4. Relationship between qPCR and histopathological findings

Fifteen fishes were positive to *S. agalactiae* and one sample was inconclusive by qPCR (Cq 39.9). This result is near to the technical

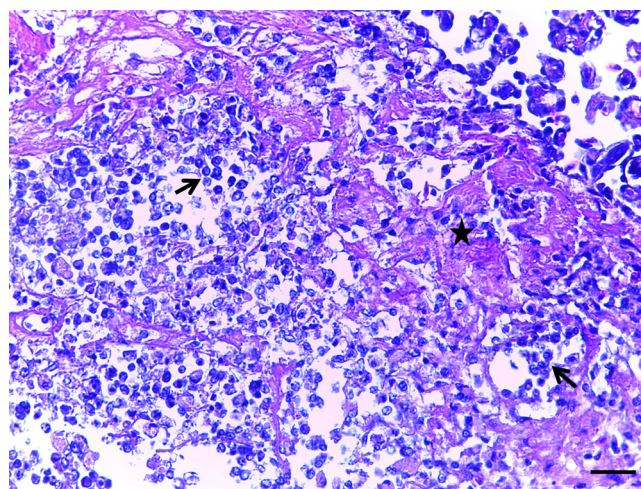


Fig. 2. Histological section of heart. Fibrinosuppurative epicarditis (star) and mononuclear infiltrate (arrow). Magnification: 400 \times . H&E.

limit of detection. The correlation between qPCR and histopathological findings (Table 2) showed fifteen fish positives (prevalence of 93.75%) with qPCR lower than 40 cycles (Cq <40). These fish presented lower Cq associated to high degree of tissue injuries.

Table 2
Values of the Cq (<40) and its correlation with the histopathological findings caused by *S. agalactiae* in *Oreochromis niloticus*.

Fish	qPCR (Cq)	Histopathological Findings							
		Brain	Eye	Gills	Heart	Muscle	Gut	Liver	Spleen
1	25.5		Panophthalmitis	Fused lamella and hyperplasic		Coagulative necrosis and mononuclear infiltrate	Necrosis and mononuclear infiltrate	Necrosis of hepatocytes Fatty and hydropic degeneration	
2	36.8	Meningitis	Ophthlmitis	Fused lamella	Epicarditis	Coagulative necrosis		Necrosis of hepatocytes and necrotic foci	Lymphocytolisis
3	37.5		Edema and ophthalmitis	Lamellar hyperplasia and fusion				Mononuclear infiltrate Fatty and hydropic degeneration	Congestion
4	36.9			Fused lamella		Coagulative necrosis and mononuclear infiltrate	Necrosis and hyperplasia	Mononuclear infiltrate	
5	38.6	Meningitis		Congestion and fused lamella	Epicarditis Myocarditis	Coagulative necrosis	Necrosis and hyperplasia	Necrosis of hepatocytes Fatty and hydropic degeneration Mononuclear infiltrate	Depletion of lymphocyte and necrosis Congestion
6	26.8	Suppurative meningitis Congestion	Panophthalmitis and dispersed pigmentary epithelium cells	Fused lamella	Chronic fibrino suppurative epicarditis Myocarditis			Necrosis of hepatocytes Fatty and hydropic degeneration Mononuclear infiltrate	
7	32.5		Panophthalmitis and congestion	Hyperplasia and congestion	Epicarditis	Coagulative necrosis		Necrosis of hepatocytes	Depletion and necrosis
8	22.0	Meningitis and congestion	Panophthalmitis	Fused lamella, congestion and hyperplasia	Chronic fibrino suppurative epicarditis Myocarditis	Coagulative necrosis	Necrosis	Necrosis of hepatocytes Fatty and hydropic degeneration.	
9	25.5	Meningitis	Ophthalmitis		Fibrino suppurative necrosis	Coagulative necrosis		Necrosis of hepatocytes	Necrosis and depletion
10	20.7	Suppurative meningitis	Ophthalmitis dispersed pigmentary epithelium cells	Lamellar hyperplasia and congestion		Coagulative necrosis		Necrosis of hepatocytes Mononuclear infiltrate	Necrosis and lymphocytolisis
11	22.5	Meningitis	Panophthalmitis	Lamellar hyperplasia and fusion	Chronic fibrino suppurative epicarditis and paniculitis Myocardial necrosis	Coagulative necrosis		Necrosis of hepatocytes Fatty and hydropic degeneration Perihepatitis fibrino suppurative.	Perisplenitis, necrosis
12	19.6		Panophthalmitis Congestion Dispersed pigmentary cells	Congestion and fused lamella	Fibrino suppurative epicarditis	Coagulative necrosis		Perihepatitis Necrotic focus Mononuclear infiltrate	Congestion Necrosis MMC increased
13	38.1	Congestion and meningitis	Ophthlmitis		Epicarditis Myocarditis	Coagulative necrosis		Fatty and hydropic degeneration Congestion Necrosis of hepatocytes	Necrosis and congestion
14	21.4		Severe panophthalmitis with dispersed pigmentary epithelium cells		Chronic fibrino suppurative epicarditis Miocarditis	Coagulative necrosis and mononuclear infiltrate		Necrosis of hepatocytes Mononuclear infiltrate	Necrosis and depletion
15	39.9					Coagulative necrosis			
16	18.3	Suppurative meningitiis		Fused lamella and congestion	Fibrino suppurative epicarditis Myocardial necrosis	Coagulative necrosis and mononuclear infiltrate		Necrosis of hepatocytes Fatty and hydropic degeneration Mononuclear infiltrate	Necrosis and lymphocytolisis

4. Discussion

The specie *S. agalactiae* is the most widely distributed in Latin American countries (Sheehan et al., 2009). However, its geographical distribution may be undervalued due to the absence of formal reporting and inadequate identification (Pulido and Iregui, 2010). In

16 sampled fishes, the molecular analysis showed fifteen positive animals. Several authors have used the PCR for the identification of *S. agalactiae*, such us multiplex PCR by detection of *Aeromonas* spp., and *Streptococcus* spp. (Lukkana et al., 2014). Also, Rodkhum et al. (2012) developed a duplex PCR for the detection of *S. agalactiae* and *S. iniae*. However, the real-time PCR assay is a faster method for

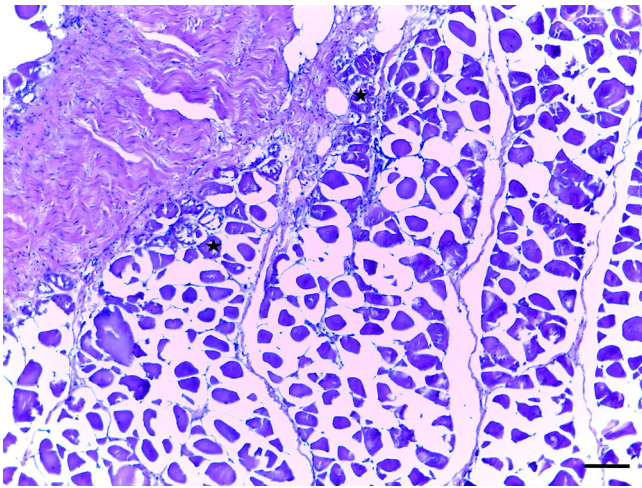


Fig. 3. Histological section of skeletal muscle. Necrosis (star). Magnification: 400 \times . H&E.

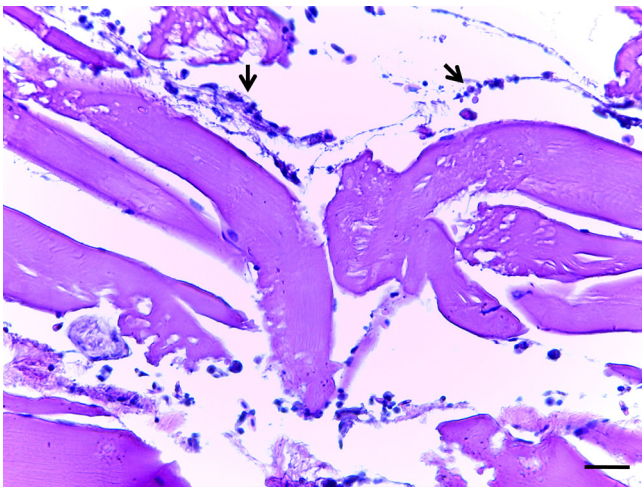


Fig. 4. Histological section of skeletal muscle. Infiltration mononuclear (arrow). Magnification: 400 \times . H&E.

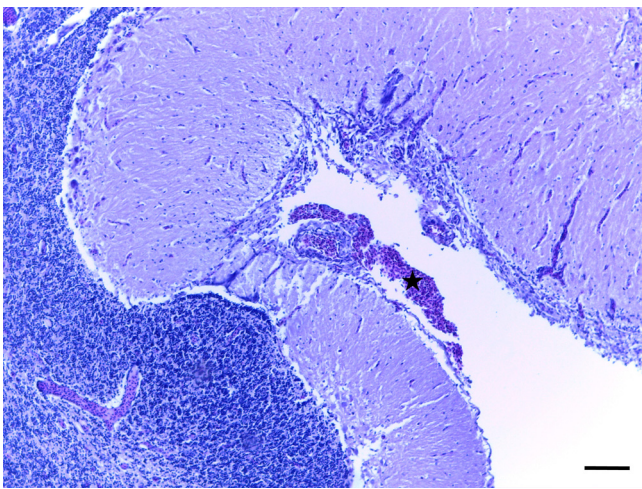


Fig. 5. Histological section of brain. Suppurative meningitis (star). Magnification: 400 \times . H&E.

identification of *S. agalactiae*, which was used in the present study. It has become the most commonly used molecular tool because it requires less labor, provides greater cost-effectiveness and is easily adapted to process large-scale samples compared with other methods (Sebastião et al., 2015). This study described the real-time PCR assay using Taqman probes that can use DNA extracted directly of a pure isolated.

Macroscopic findings found at the necropsy, such as unilateral or bilateral exophthalmia, hyphema, corneal opacity, ascites, melanosis, bleeding in the anal pore, haemorrhagic and necrotic lesions on the skin and muscle tissue, fluid in the abdominal cavity, hepatomegaly, splenomegaly are suggestive of *Streptococcus agalactiae* infection, in accordance with the described by Amal and Zamri-Saad (2011), and also reported by Salvador et al. (2005) and Figueiredo et al. (2006) in tilapias infected naturally.

The histopathological description of the fish sampled showed correlation with qPCR results, because animals with low values of Cq (<40) had more histopathological lesions. According with Iregui et al. (2014), to identify DNA in tissues do not mean that the fish has the disease, and only the correlation between the results from molecular techniques, anamnesis and microscopic findings can offer more reliable diagnoses (Sebastião et al., 2015). In this study, the tissues most affected by the positive fish were the heart, liver, eye, spleen and brain. Such results suggest that they would be target organs for this bacterium, which has also been reported by Alsaïd et al. (2013). These findings indicate septic process and spread this agent as reported by Yuasa et al. (2008). Also, the findings of epicarditis and meningitis found in this study are consistent with those described by Zamri-Saad et al. (2010) in *O. niloticus* infected naturally with *S. agalactiae*. Alsaïd et al. (2013) and Pulido et al. (2004) mentioned that inflammation in the epicardium may vary in different degrees, while other organs could be less affected, corroborating the findings observed in this study with types of acute or chronic epicarditis. In the spleen was found congestion, necrotic foci and increased of MMC. Similar histological lesions in the splenic tissue of red tilapia were described by Alsaïd et al. (2013). Other authors observed increased number of MMC in the spleen of fish during bacterial infection (Manrique et al., 2014).

The lesions observed in the liver, as necrosis and mononuclear infiltration, described in this study are consistent with those reported by Mohamed et al. (2014) in tilapias infected experimentally with *Streptococcus* spp. It is noteworthy that the infiltration of mononuclear cells, mainly macrophages, observed in this study in all organs were also described by Filho et al. (2009) and Iregui et al. (2014).

In Asia and Latin American, were isolated near of 500 cases of *Streptococcus* sp. coming from tilapia, which 82% of prevalence were isolates of *S. agalactiae* and 18% of *S. iniae* in the *Oreochromis* sp. Some authors mention that *S. agalactiae* is among the most prevalent species in Latin American countries. In Brazil and Colombia, this bacterial infection has caused high mortality in tilapia farms (Iregui et al., 2014; Marcusso et al., 2015). This study was the first to identify by qPCR *S. agalactiae* in naturally infected tilapia from nursery and grow-out phases in Piura, Peru, and to characterize histopathological lesions caused by this bacterium. The correlation between the histopathological findings with the qPCR can provide greater support in relation to the diagnostic and specific treatment for this disease.

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