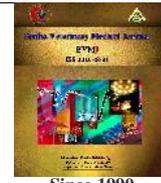




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Control of some septicemic bacterial infections in *Oreochromis niloticus*, with special reference to their antimicrobial resistant genes

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ABSTRACT

The current study aimed to control the bacterial infection associated with summer mortalities in Nile tilapia. The sensitivity of the isolated bacteria, including *Aeromonas Veronii* (*A. veronii*) strain (*HY1*, *HY2*, *HY3*, *HY4* and *HY6*), *Citrobacter freundii* (*C. freundii*), *Proteus vulgaris* (*P. vulgaris*) and *pseudomonas fluorescens* (*P. fluorescens*) for antibiotic was tested and antibiotic resistant genes were investigated. In order to verify the result of antibiotic sensitivity; experimental trial was performed to determine the effect of florfenicol and erythromycin against Nile tilapia (*Oreochromis niloticus*) infection with different pathogenic bacteria [4 strains of *A. veronii* (*AHY2*), *AHY3*), *AHY4*) and *AHY6*), *C. freundii*, *P. fluorescens* and *P. vulgaris*]. The results revealed that all tested bacterial isolates were 100 % sensitive to florfenicol and erythromycin, 50% sensitive to gentamycin, 25 % sensitive to nalidixic acid and sulphamethoxazol-trimethoprim, while all bacterial isolates were 100% resistant to oxytetracycline and tetracycline lincomycin, ofloxacin and penicillin, 87.5% resistant to ampicillin, 75% resistant to sulphamethoxazol-trimethoprim, 62.5% for nalidixic acid and 37.5% for gentamycin. The relative percent survival (RPS) of challenged medicated fish with erythromycin ranged from 45.0-91.7%. Where, the survival % in *HY3* infected groups was the highest (91.7%) followed by *C. freundii* (87.5%), *HY6* (83.3 %) and *P. vulgaris* (81.3%).

1. INTRODUCTION

Tilapia culture is stepping toward the top in the field of aquaculture industry, where the world tilapia aquaculture production is grown 12% per year (FAO, 2017). Tilapia fish farms are vulnerable to various bacterial infections which necessitate the usage of antimicrobials. However, with the expansion of fish farms and intensification systems, the overuse of antimicrobials leads to emerging of new bacterial strains which acquired resistance to the antimicrobial agents (Watts et al., 2017). Most of antimicrobial resistance (AMR) occurs through mutation, horizontal gene transfer (HGT), or through natural transformation, transduction, or conjugation (von Wintersdorff et al., 2017). The incidence of antimicrobial resistance in fish shows the chance of transfer of AMR genes to humans, and the difficulty in controlling diseases (Santos and Ramos, 2018), which lead to their limitation and developing laws and legislations for antimicrobial use in aquaculture (Okocha et al., 2018). In order to reduce AMR, a necessary attempt to determine the bacterial sensitivity to the approved antimicrobials should be adopted (AVMA 2006). Limitation are facing the farmers for choosing the antimicrobial agent, hence; only three antimicrobials; oxytetracycline (Terramycin® 200), florfenicol (Aquaflor®) and ormetoprim and sulfadimethoxine (Romet30) are approved by the United States Food and Drug Administration (USFDA) as feed additives in aquaculture sector (Manyi-Loh et al., 2018) Florfenicol has been used as medicated feed for controlling *Vibrio anguillarum*,

Aeromonas salmonicida, *Streptococcus iniae*, *Listonella anguillarum*, and *Edwardsiella ictaluri* in different fish species (Darwish, 2007., Gaunt et al., 2003., Samuelsen et al., 1998., Samuelsen and Bergh, 2004., Seljestokken et al., 2006). Although erythromycin is not found in the approved antimicrobials list, but it is proved effective in treatment *Flavobacterium psychrophilum* (Jarau et al., 2019), and it was used as a prophylactic against bacterial kidney disease (Elliot, 2017).

The aim of this work was to control the bacterial infection based on the data obtained from the antimicrobial profile and the AMR genes

2. MATERIAL AND METHODS

2.1. Bacterial isolates

In the present study; *A. veronii* (*HY1*, *HY2*, *HY3*, *HY4* and *HY6*), *C. freundii*, *P. vulgaris* and *P. fluorescens* isolated from diseased Nile tilapia during summer mass mortality were used. All the isolates were previously identified on phenotypic and molecular basis (Youssuf et al., 2020).

2.2. Antimicrobial sensitivity test and antimicrobial resistance index (AMR)

The selected bacterial strains were grown overnight in Muller Hinton broth (MHB) (Oxoid, UK) at 28°C for 24 hr. Then the bacterial suspension was inoculated on Muller Hinton Agar (MHA) (Oxoid, UK) plates and the discs of 11 chemotherapeutic agents (Oxoid, UK) were placed firmly on the surface of the agar. The plates were incubated at 28°C

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for 24 h. The results were determined by measuring the diameter of the inhibition zones and the interpretation of inhibition zones were carried out using NCCLS (2000) guidelines (Table 1). The multiple antimicrobial resistant (MAR) index was calculated according to Schwarz et al., (2010) by dividing the total number of resistance to antimicrobials by each isolate on the total numbers of tested antimicrobials.

2.3. Determination of antimicrobial resistant genes of the pathogenic bacteria

Based on the antibiogram result, a total of seven pathogenic strains including four strains of *A. veronii* (*AHY2*, *AHY3*, *AHY4*) and *AHY6*), *C. freundii*, *P. flurosence* and *P. vulgaris* were selected for determination of antimicrobial resistant genes. DNA extraction was performed from strains resistant to more than one antibiotic, using QIAamp DNA mini kit according to the manufacturer instructions. Identification of different antimicrobial resistant genes coding was performed using PCR amplification with specific forward and reverse oligonucleotide primers listed in table (2). PCR reactions were carried out in 25 µl volumes containing 6 µl of DNA, 1 µl of each primer (20 pmol), and 12.5 µl of Emerald Amp GT PCR Master Mix (2x premix) and 4.5 µl of PCR grade water. All PCR reactions were done at 94°C for 5 min, 35cycles at 94°C for 30 s, 60°C for 1 min, 72°C for 1 min, followed by a final elongation at 72 °C for 10 min. The obtained PCR products were analyzed by electrophoresis.

2.4. Using chemotherapy for controlling of pathogenic bacterial infection in Nile tilapia

2.4.1. Preparation of medicated diet

Florfenicol (Floribiotic 100 %, ATCO Pharma) medicated diet was prepared at concentration of 15 mg/kg fish following Soto *et al.*, (2010), while, erythromycin (Ucimycin P.F, 20%, Muveco, veterinary pharmaceuticals, fodder concentrates company) medicated diet was carried out at a concentration of 50 mg/ kg fish body weight according to Phuoc *et al.* (2010). The required dose of the chemotherapy was dissolved in an appropriate amount of tap water and mixed with 1000 g crushed commercial diet to produce dough. The diet was re-pelleted using hand pelletizer, dried at room temperature for 48 hr. The medicated diets were packaged and stored at 4°C.

2.4.2. Experimental Design

About 440 Nile tilapia fingerlings (20 ± 5 g average body weight) were transported alive from fish farm at Kafr El-Sheikh to the wet laboratory of Aquatic Animal Diseases and Management, Faculty of Veterinary Medicine, Moshtohor, Benha University. Fish were kept in well prepared fiber glass tanks (750 L capacity) for 2 weeks to be acclimated for laboratory conditions. Each tank was supplied with an air pump and dechlorinated tap water. Then fish were maintained at stocking densities of 10 fish per glass aquaria (100 × 100 × 50), the water parameters were adjusted as following; water temperature 27±2 °C, dissolved oxygen 6±0.5mg/L, total ammonia concentration 0.53±0.07 mg/L and pH 7±0.2. Fish were fed with 4% body weight basal diet (30% protein) twice daily (Mesko, 1985). After two weeks acclimation, fish were randomly distributed into 22 group in duplicate (10 fish/replicate) in well prepared glass aquaria supplied with aeration and de-chlorinated tap water. The health status of fish was determined during acclimation period according to Austin and Austin (2007).

Table 1 The different antimicrobial discs used in sensitivity test with interpretation of the inhibitory zone

Antibiotic	Disc content (µg)	Diameter of the inhibiting zone around the disc mm		
		Resistant	Intermediate	Sensitive
Florfenicol	30 µg	14	15-18	19
Erythromycin	15 µg	13	14-22	23
Penicillin G	10 U	19	20-27	28
Ampicillin	10 µg	15	16-18	19
Lincomycin	10 µg	22	23-25	26
Ofloxacin	10 µg	15	16-21	22
trimethoprim-sulphamethoxazol	25 µg	22	23-25	26
Gentamycin	10 µg	12	13-14	15
nalidixic acid	30 µg	22	23-25	26
Tetracycline	30 µg	22	23-25	26
Oxytetracycline	30 µg	22	23-25	26

Table 2. Oligonucleotide primers sequences Metabion (Germany) which used in determination of antimicrobial resistant genes

Primer	Sequence	Amplified product	Reference
<i>blaTEM</i>	F/ATCAGCAATAAACCAGC	516 bp	Colom <i>et al.</i> , 2003
	R/CCCCGAAGAACGTTTTTC		
<i>QnrS</i>	F/ACGACATTCGTCAACTGCAA	417 bp	Robicsek <i>et al.</i> , 2006
	R/TAAATTGGCACCCCTGTAGGC		
<i>SulI</i>	F/CGGCGTGGGTACCTGAACG	433 bp	Ibekwe <i>et al.</i> , 2011
	R/GCCGATCGCGTGAAGTTCCG		
<i>tetA(A)</i>	F/GGTTCACCTCGAACGACGTCA	576 bp	Randall <i>et al.</i> 2004
	R/CTGTCCGACAAGTTGCATGA		
<i>MphA</i>	F/GTGAGGAGGAGCTTCGCGAG	403 bp	Nguyen <i>et al.</i> , 2009
	R/TGCCGACGACTCGGAGGTC		
<i>DfrA</i>	F/TGGTAGCTATATCGAAGAATGGAGT	425 bp	Grape <i>et al.</i> , 2007
	R/TATGTTAGAGCGAAGTCTGGGTA		
<i>ErmB</i>	F/GAAAAAGTACTCAACCAAATA	639 bp	Nguyen <i>et al.</i> , 2009
	R/AATTTAAGTACCGTTACT		

blaTEM (penicillin), *QnrS*(quinolones), *SulI* (sulphonamide), *tetA(A)* (tetracyclines), *MphAI* (macrolides), *DfrAI* (trimethoprim) and *ErmB* (lincomycin).

Fish were divided into three main groups, main group1; MG1; (with 7 subgroups) in which fish were fed florfenicol medicated diet and fish in each subgroup was (IP) challenged with 0.2 ml of the identified pathogenic bacteria (4 strains of *A. veronii* (AHY2), *A* (HY3), *A*(HY4) and *A*(HY6)), *C. freundii*, *P. flurosence* and *p. vulgaris*) to achieve 1.8×10^8 cells/fish. Main group2; MG2; (with 7 subgroups) in which fish were fed erythromycin medicated diet and fish in each subgroup was (IP) challenged with 0.2 ml of the same identified pathogenic bacteria (4 strains of *A. veronii* [(AHY2), AHY3), AHY4) and AHY6]), *C. freundii*, *P. flurosence* and *p. vulgaris*) to achieve 1.8×10^8 cells/fish. Main group3; MG3; (with 7 subgroups) in which fish were fed basal diet and (IP) challenged with 0.2 ml of the pathogenic bacteria [4 strains of *A. veronii* (AHY2), AHY3), AHY4) and AHY6]), *C. freundii*, *P. flurosence* and *p. vulgaris*) to achieve 1.8×10^8 cells/fish, these groups were considered as control positive groups. Control negative (non-infected group) was fed basal diet. The treatment in all groups was started just after appearance of septicemic signs. Fish in experimentally treated and control groups was fed at 4% of their body weight twice a day along the experimental period (10 days). Clinical signs, post-mortem lesions and mortalities were recorded during the experiment. The protocol of work was approved with a number (BUFVTM-

06-07-20), and monitored by the Ethics Committee of the Faculty of Veterinary Medicine, Benha University, Egypt.

3. RESULTS

The antibiogram profile revealed that the (AHY2) strain showed resistance to ampicillin, lincomycin, tetracycline, ofloxacin, oxytetracycline and penicillin. (AHY3) strain and *C. freundii* showed resistance to ampicillin, lincomycin, nalidixic acid, tetracycline, ofloxacin, oxytetracycline, penicillin and sulphamethoxazol- trimethoprim. (AHY4) and *P. flurosence* showed resistance to ampicillin, lincomycin, nalidixic acid, tetracycline, ofloxacin, oxytetracycline, penicillin, sulphamethoxazol- trimethoprim and gentamycin. (AHY6) strain) showed resistance to lincomycin, nalidixic acid, tetracycline, ofloxacin, oxytetracycline, penicillin, Sulphamethoxazol- trimethoprim and gentamycin and (*P. vulgaris*) showed resistance to ampicillin, lincomycin, tetracycline, ofloxacin, oxytetracycline, penicillin and sulphamethoxazol- trimethoprim. Also strain (AHY1) showed moderate resistance to the six chemotherapeutic agents (ampicillin, lincomycin, tetracycline, ofloxacin, oxytetracycline and penicillin) at rate of 54.5% (Table 3).

Table (3): Degree of sensitivity of bacterial isolates to every antimicrobial agent.

Bacterial Isolates	Antimicrobial agents										
	ampicillin (Amp) 10 µg	gentamycin (GN) 10 µg	lincomycin (MY) 10 µg	nalidixic acid (NA) 30 µg	tetracycline (TE) 30 µg	ofloxacin (OFX) 10 µg	oxytetracycline (OT) 30 µg	sulphamethoxazol-trimethoprim (SXT) 25 µg	Penicillin G 10U	Erythromycin 15 µg	Florfenicol 30 µg
<i>A. veronii</i> (HY1)	R	S	R	S	R	R	R	S	R	S	S
<i>A. veronii</i> (HY2)	R	S	R	S	R	R	R	S	R	S	S
<i>A. veronii</i> (HY3)	R	S	R	R	R	R	R	R	R	S	S
<i>A. veronii</i> (HY4)	R	R	R	R	R	R	R	R	R	S	S
<i>A. veronii</i> (HY6)	I	R	R	R	R	R	R	R	R	S	S
<i>Proteus vulgaris</i>	R	I	R	I	R	R	R	R	R	S	S
<i>Pseudomonas Flurosence</i>	R	R	R	R	R	R	R	R	R	S	S
<i>Citrobacter freundii</i>	R	S	R	R	R	R	R	R	R	S	S
Degree of sensitivity of bacterial isolates to every antimicrobial agent											
% of R	87.5	37.5	100	62.5	100	100	100	75	100	0	0
% of I	12.5	12.5	0	12.5	0	0	0	0	0	0	0
% of S	0	50	0	25	0	0	0	25	0	100	100

S: Sensitive. R: Resistant. I: Intermediate. %= No. of R or I or S bacteria × 100 Total No. of bacteria

Data of the multiple antimicrobial resistances (MAR) index are present in Table 4. Data showed that the multidrug resistance pattern of the bacterial isolates and MAR index values ranged between 0.54 and 0.81.

All tested bacterial isolates [*A. veronii* (HY1, HY2, HY3, HY4 and HY6), *C. freundii*, *P. vulgaris* and *P. flurosence*] were 100 % sensitive to florfenicol and erythromycin, 50% sensitive to gentamycin, 25 % sensitive to nalidixic acid and sulphamethoxazol-trimethoprim, while sensitivity to other antibiotic discs was 0%. In addition, all bacterial isolates were 100% resistant to oxytetracycline and tetracycline lincomycin, ofloxacin and penicillin, 87.5% resistant to ampicillin, 75% resistant to sulphamethoxazol-trimethoprim, 62.5% for nalidixic acid and 37.5% for gentamycin (Table 4).

Table 4. Percentages of antimicrobial resistance to the different antimicrobials used in the antimicrobial sensitivity test

Aeromonas spp. Isolates	Antimicrobial profile	
	No. of R. agents	MAR Index
<i>A</i> (HY1)	6	0.54
<i>A</i> (HY2)	6	0.54
<i>A</i> (HY3)	8	0.72
<i>A</i> (HY4)	9	0.81
<i>A</i> (HY6)	8	0.72
<i>Proteus vulgaris</i>	8	0.72
<i>P. flurosence</i>	9	0.81
<i>Citrobacter freundii</i>	8	0.72

The antimicrobial resistant genes of [(*A. HY2*), (*A. HY3*), (*A. HY4*), (*A. HY6*)], *P. vulgaris*, *P. fluorescence* and *C. freundii*) were tested. These isolates were found to be associated with all virulent genes (Ermb (lincomycin), Tem (penicillin), Teta

(tetracyclines), Sul1 (sulphonamide), Dfra (trimethoprim), Mpha (macrolides) and Qnr (quinolones) related to these antibiotics (Fig. 1 and Fig. 2).

Moreover, they were resistant to frequently used antibiotics for human infections. The bacterial strains revealed resistance to penicillin at 516 bp, Trimethoprim at 425 bp, macrolides at 403 bp, quinolones at 417 bp, tetracycline at 576 bp, lincomycin at 639 bp and sulphonamide at 433 bp. The treatment trial indicated that the Relative percent survival (RPS) of challenged medicated fish with florfenicol ranged from 44-100% within ten days post-infection, being the highest (100%), especially with *Aeromonas* strains HY3 (83.3%) and HY6. However, 100% survivability was evident for *Proteus vulgaris*, *C. freundii* and *P. fluorescense* (Table 5).

Relative percent survival (RPS) of challenged medicated fish with erythromycin ranged from 39-91.7% in *A. veronii* strains, where the highest survival of 83.3 and 91.7% was evident in HY6 and HY3; respectively. Similarly, higher survivability was noticed in *C. freundii* (87.5%), HY6 (83.3%) and *P. vulgaris* (81.3%) (Table 6). The challenged-medicated fish showed no abnormal signs and appeared healthy. The challenged control group survival rate was 0%

with aeromonase strain (A HY2), 10% for (A HY4), 40% for A HY3, AHY6), 20% for (*P. vulgaris* and *C. freundii*) and 60% for *P. fluorescense*. Meanwhile, *Aeromonas* strains (HY2 and HY4) reached 50% mortality.

4. DISCUSSION

Bacterial diseases are one of the many challenges facing aquaculture development (Aly, 2013). Among others, motile *Aeromonas* septicemia (MAS), caused by *Aeromonas* spp., is one of the important bacterial infections, leading to mass mortalities in fish and severe economic losses (Noga, 2010, Assefa and Abunna, 2018). Antibiotic usage in fish farms is widespread either for the prevention or treatment of bacterial infections in fish (Cabello et al., 2013), however; their misuse has led to the emergence of bacterial species with multiple drug resistance all over the world. Our data revealed that most of bacterial isolates showed high resistance rates to several antimicrobials, since strains showed resistance to at least 6 out of 11 evaluated antibiotics.

Table 5. Efficacy of florfenicol in the treatment of experimentally *O. niloticus* challenged with the pathogenic bacterial isolates.

Bacterial isolates	(Control – ve) % of dead fish	(Control + ve) % of dead fish	(Treated group) % of dead fish	Relative percent survival (RPS) of treated group
A(HY2)	0%	100%	50%	50
A(HY3)	0%	60%	0%	100
A(HY4)	0%	90%	50%	44.4
A(HY6)	0%	60%	10%	83.3
<i>Proteus vulgaris</i>	0%	80%	0%	100
<i>P. fluorescense</i>	0%	40%	0%	100
<i>C. freundii</i>	0%	80%	0%	100

Table 6. Efficacy of erythromycin in the treatment of experimentally *O. niloticus* challenged with the pathogenic bacterial isolates.

Bacterial isolates	(Control – ve) % of dead fish	(Control + ve) % of dead fish	(Treated group) % of dead fish	Relative percent survival (RPS) of treated group
A(HY2)	0%	100%	55%	45
A(HY3)	0%	60%	5 %	91.7
A(HY4)	0%	90%	55 %	38.9
A(HY6)	0%	60%	10 %	83.3
<i>Proteus vulgaris</i>	0%	80%	15 %	81.3
<i>P. fluorescense</i>	0%	40%	20 %	50
<i>C. freundii</i>	0%	80%	10%	87.5

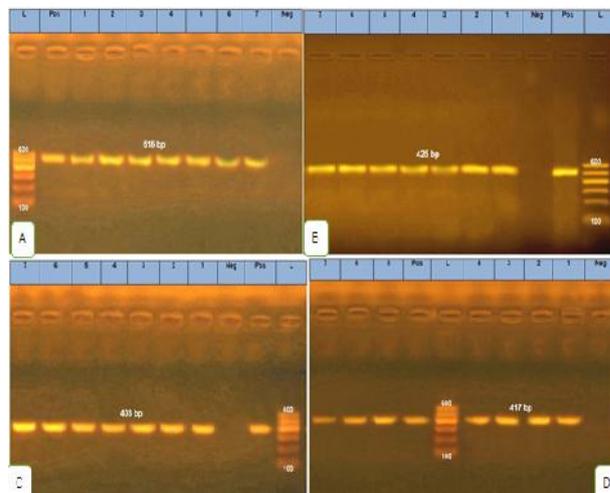


Fig. 1 PCR amplification of A): blaTEM gene related to (penicillin) at 516 bp. B): dfra gene related to (Trimethoprim) at 425 bp. C): mpha gene related to (macrolides) at 403 bp. D) qnrA gene related to (quinolones) at 417 bp. Lane: 1-7 (A(HY2), A(HY3), A(HY4),

A(HY6), Proteus vulgaris, Pseudomonas fluorescense and Citrobacter freundii) positive. Neg=Negative control, pos=Positive control, L=Marker

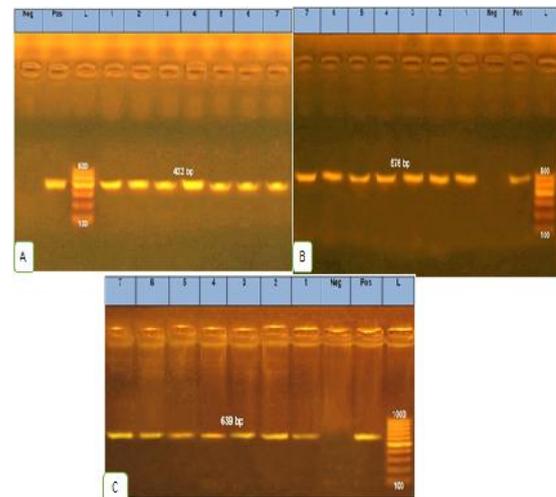


Fig. 2 PCR amplification of A): sulI gene related to (sulphonamide) at 433. B) tetA gene related to (tetracyclin) at 576. C): ermB gene related to (lincomycin) at 639 bp. Lane: 1-

7 (A(HY2), A(HY3), A(HY4), A(HY6), *Proteus vulgaris*, *Pseudomonas fluorescense* and *Citrobacter freundii*) positive. Neg=Negative control, pos=Positive control, L=Marker.

All bacterial isolates were 100% resistant to oxytetracycline, tetracycline, lincomycin, ofloxacin and penicillin. This result was consistent with previous studies (Rhodes et al. (2000), Kim et al. (2004) and Hatha et al. (2005)). The modest extent of resistance in our study is surprising as tetracyclines are widely used for control of most bacterial fish diseases in aquaculture in Egypt. A considerable overall increase in drug resistance of fish pathogenic bacteria is parallel with the extensive use of chemotherapeutic agents, which has created a great difficulty in the treatment of bacterial infections in fish (Aoki et al., 1992). It was found that all tested bacterial isolates were highly sensitive to florfenicol (100%). Similarly, Abdel-Latif and Sedeek, (2017) found that isolated strains about (144 Enterobacteriaceae strains) which were derived from diseased fish and were markedly sensitive to gentamicin, florfenicol, and flumequine, moderately sensitive to doxycycline and less sensitive to oxytetracycline, erythromycin, and ampicillin. But the difference in sensitivity to erythromycin in our study may be due to difference in bacterial isolates and extensive use and misuse of these antibiotics. Our results showed resistance to more than one class of antibiotic (Multiple drug resistance) which has been reported for fish pathogens in several studies (Schmidt et al., 2000 and Hatha et al., 2005). The high resistance levels of bacteria to antimicrobials, which due to the frequent use of antibiotics either as growth promoters or therapeutic agents for various diseases. MAR index of our isolates showed high ranges between 0.54 and 0.81. In the same context, MAR index ranged between 0.27 and 0.82 in the study of El Gohary et al., (2020).

A variety of resistance patterns was observed in our study which explained by determination of antimicrobial resistant genes as the isolates (AHY2), AHY3), AHY4), AHY6), *P. vulgaris*, *P. fluorescense* and *C. freundii* were found to be associated with all virulence genes detected; ErmB (lincomycin), Tem (penicillin), Teta (tetracyclines), SulI (sulphonamide), Dfra (trimethoprim), Mpha (macrolides) and Qnr (quinolones). The spread of tet genes is often facilitated by their location on mobile genetic elements, such as plasmids and transposons (Chopra and Roberts, 2001). Plasmid-mediated quinolone resistance in bacteria associated with fish farms has been detected in several countries. The identification of QnrS determinants in *Aeromonas* spp. suggests that they may act as an environmental reservoir of qnrS genes, as already described for tet genes (Rhodes et al., 2000 and Schmidt et al., 2001). Some of our bacterial isolates confirmed to have resistant genes to some antibiotics, whereas *in vitro* they proved to be sensitive to those antibiotics, as (A(HY2) was sensitive to sulphamethoxazol-trimethoprim, while A(HY4) and A(HY2) were sensitive to lincomycin. Our explanation is those strains may have nonfunctional genes which doesn't work and this came in accordance to Zheng et al., (2007) who reported that Pseudogenes are DNA fragments, related to recognized genes, which have lost their protein-coding ability or are no longer expressed in the cell. Pseudogenes originate from retro transposition or genomic duplication of functional genes and become "genomic fossils" that are non-functional attributed to mutations that inhibit gene transcription, such as in the region of the promoter of genes, or fatally alter the translation of the gene, such as premature stop codons or frameshifts. Pseudogenes resulting from the retro transposition of an RNA intermediate are known as processed pseudogenes; pseudogenes that arise from the

genomic remains of duplicated genes or residues of inactivated genes are non-processed pseudogenes (Zheng et al., 2007). Additionally, David et al., (2009) about his work on *E. coli*, stated that although a wide variety of expression hosts would need to be employed, for the detection of cloned functional resistance from different bacterial genera. The transformation of the transmitted genes into functional genes in new hosts would require comprehensive mutation and recombination "tailoring." Native proponents need to be changed to allow transcription and translation in the new hosts.

The use of feed treatments is the most convenient method on the large scale aquaculture, to avoid the stress and reduce labor (Robb and Crampton, 2013), but it is important to start feeding before fish stop eating (i.e; tend to be sick); otherwise the benefit of the delivered medication wouldn't be achieved.

The experimental trial indicated that survivability of challenged medicated fish with florfenicol was nearly at the same erythromycin treated fish, compared to the infected non-treated groups, which indicate the effectiveness in controlling the infection. Similarly, Darwish, (2007) reported that the daily florfenicol dose of 15 mg/kg BW for 10 d against *Streptococcus iniae* infection in Sunshine Bass significantly increased the final survival % to a level that was statistically indistinguishable from that of the negative control group (100% survival). Additionally, Nile tilapia challenged with *Enterococcus faecalis* and fed florfenicol medicated diet (15 mg/kg fish body weight) showed higher survivability (83.3%) than control group (66.7 %) (Abd El-latif et al., 2013).

5. CONCLUSIONS

It could be concluded that both erythromycin and florfenicol proved effective in treating bacterial infections associated with summer mortality syndrome. However, our results revealed that many important bacteria isolated from Nile tilapia farms have multiple drug resistance and this is due to the misuse of antimicrobial agents. So, we have to find another alternative way to control bacterial diseases in tilapia culture.

CONFLICT OF INTEREST

The authors declare that they don't have any conflict of interest.

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