

Yersiniosis in *Oreochromis niloticus*; Prevalence, Antimicrobial Resistance and Immunological Response to Ascorbic Acid and Difloxacin

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ABSTRACT

Yersiniosis is a significant disease of economic importance in salmonids and non-salmonid fishes. However, few studies have investigated the pathogenesis, preventive, and treatment measures for this bacterial pathogen. Thus, the pathogenesis of *Yersinia ruckeri* in cultured *Oreochromis niloticus* in different localities was investigated throughout 2019. Besides, two experiments were conducted, the first feeding experiment based upon dietary supplementation by Ascorbic Acid and Difloxacin. In the second experiment, the supplemented fish groups were challenged by pathogenic *Y. ruckeri* strains, treated by Difloxacin, then the hematological, and immunological assays were analyzed. The supplemented fish groups by Ascorbic Acid and Difloxacin showed an increase in serum lysozyme, NO and SOD concentrations, phagocytic activity, and Interleukin IL-2 and IL-10 levels. The challenged fish by *Y. ruckeri* reported significant disturbances in transferases activities (ALT and AST), urea, total protein, and albumin concentrations. Moreover, RBC, Hb, PCV, nitroblue tetrazolium, lysozyme, NO and SOD concentrations, and Phagocytic activity were declined, with elevated interleukins level compared to the control. Thus, Ascorbic Acid supplementation and Difloxacin dosing could enhance the immune response, improve health status, and curtail the increased emergence of antibiotic-resistant bacteria.

INTRODUCTION

Nile Tilapia (*Oreochromis niloticus*) is of the most commercial fish species in Egypt and worldwide (Suresh, 2003). Despite the obstacles facing its production, but it is still considered the best native source of animal protein, which contributes to bridging the gap per capita. Disease outbreaks have been a major limitation to the dilation of aquaculture which has a significant action on the economic development in many countries (Aly, 2009; Abdel-Aziz et al., 2020; Van Doan et al., 2020; Xia et al., 2020).

Enterobacteriaceae species; *Edwardsiella tarda*, *E. coli*, *Yersinia ruckeri*, and *Proteus retgerri* are bacterial microorganisms that cause severe economic losses among fish farms worldwide. Enteric redmouth disease (ERM) or Yersiniosis, caused by *Yersinia ruckeri*, Gram-negative bacterium that leads to significant economic losses in different fish species (Ross *et al.*, 1966; Horne and Barnes, 1999). This disease was first reported in rainbow trout *Oncorhynchus mykiss* in the United States in 1958, then become endemic in North America (Bullock *et al.*, 1977; Daligault *et al.*, 2014; Navas *et al.*, 2014).

Other marine fish species such as turbot *Scophthalmus maximus*, seabass *Dicentrarchus labrax*, and seabream *Sparus auratus* have also exhibited the infection (Bullock and Cipriano, 1990). Recently, it is reported throughout North and South America, Europe, Australia, South Africa, the Middle East, and China (Tobback *et al.*, 2007; Shaowu *et al.*, 2013). In Egypt, *Y. ruckeri* infection has been recorded in *O. niloticus* (Hussein *et al.*, 1997) and common carp and African catfish (Abd-El Latief, 2001).

The specific and non-specific immune restraints to natural or experimental infection by *Y. ruckeri* strains have been established (Anderson *et al.*, 1979; Cossarini-Dunier, 1986; Salah *et al.*, 2012). Following the immersion of rainbow trout by *Y. ruckeri*, high numbers of *Y. ruckeri* were attached to the gill mucus, and it was spotted within the gill capillaries, leading to septicemia and colonization of the internal organs (Tobback *et al.*, 2009). Clinically, Yersiniosis is characterized by the development of acute and/or chronic entero-septicemia, exophthalmia, and hemorrhage on the mouth, skin, gills, and intestine (Fuhrmann *et al.*, 1983; Rigos and Stevenson, 2001). The infection could be seen in a per-acute or acute form in young fish stages (Danley *et al.*, 1999; Rigos and Stevenson, 2001).

Nutritional supplementation could influence the immune response of fish, play a critical role in improving health status, and subsequently provide the ability to resist diseases (Blazer, 1992; Lall, 2000; Rahimi *et al.*, 2015). They include essential nutrients such as proteins, fatty acids, vitamins, polysaccharides, and some minerals that have vital importance to promote their immune response (Barrows *et al.*, 2008). Vitamins C and E are among the most important nutrients which influencing the immune performance of fish (Shiau and Hsu, 2002; Puangkaew *et al.*, 2004). Vitamin C is a water-soluble antioxidant and an essential microelement in biological fluids associated with biochemical reactions in cells and tissues.

Besides, antibiotics could be used in aquaculture as therapeutic and prophylactic agents (Aly, 2009; Pham *et al.*, 2015). But, the extensive use and misuse of antibiotics as prophylactic agents share in the development and spread of antibiotic-resistant microorganisms in the environment, among human, and animal food (Cabello, 2006). Although some antimicrobials can protect against microbial infection, public health and environmental hazards must be taken into consideration.

Thus two aims were planned in this study, the first was to investigate the effect of dietary supplementation of Ascorbic acid and prophylactic antibiotics to the healthy fish and study the pathogenesis of *Y. ruckeri* in cultured *O. niloticus* in different localities. The second aim was to elucidate a treatment strategy by the antibiotic following the experimental infection of Nile

tilapia with pathogenic *Y. ruckeri*. Consequently, the hematological, serum biochemical, and immunological parameters were assessed.

MATERIAL AND METHODS

Fish for studying the pathogenesis of natural infection

Fish sampling was performed according to the legislation of animal welfare following the guideline addressed for on humane treatment of fish (Adams *et al.*, 2002). A total of 100 cultured *O. niloticus* with an average weight of 100 ± 25 gm were collected from five earthen pods farms (Farm1- Farm 5), in Sharkia Governorate to study the pathogenesis of *Y. ruckeri*. Then the fish were transported immediately alive in sterile polyethylene bags to the Pathology Laboratory, Faculty of Veterinary Medicine, Suez Canal University.

Bacteriological examination

The bacteriological isolates were gathered from the body surface, gills, and intestine and cultured aseptically on Tryptic Soy Broth. Following incubation at 25 °C for 48h, they streaked onto Heart Infusion Agar BHIA or Tryptic soy TSA media and incubated again at 25 °C for 48h. Then different pure colonies were selected and sub-cultured on selective medium and Yeast Extract Glucose Agar (Michel *et al.*, 1997) *Y. ruckeri* Selective Medium (Furones *et al.*, 1993). The morphological and biochemical characterization was performed according to Waltman and Shotts, (1984), Furones *et al.* (1993), Elmeer *et al.* (1997). More confirmatory biochemical tests were performed using a commercial API 20 E system.

Determination of antimicrobial resistance

It was determined by disk diffusion on The Mueller–Hinton agar (Difco.) using six antimicrobial agents (Chloramphenicol (C 30 µg), Sulphamethoxazole & Trimethoprim (SXT 25µg), Difloxacin (D10µg), Ampicillin (AMP 10µg), Oxytetracycline (OT 30µg), Erythromycin (E 10 mg) (Oxoid). The concentration of antimicrobials in the disks and the inhibition zone breakpoint values of the resistance were interpreted and the disk diffusion assays were registered according to the recommendations of the CLSI (2005 a & b).

Experiment 1: Dietary supplementation by Ascorbic acid and Difloxacin (1-30 day)

A total of 360 healthy *O. niloticus* of 70 ± 12 g body weight were assembled from a commercial farm. The fish were transported directly to the wet laboratory, then distributed in circular fiber-glassed fiber each of 1000 L capacity. All the requirements of the closed recirculating and filtration system, the water quality, feeding regime, and the health status of fish were managed.

Following acclimatization, the fish were divided into three triplicated groups: G1; control group fed basal diet only. G2; fed basal diet supplemented with ascorbic acid at 1000 mg/kg b.wt. (Blazer, 1992) for 30 days. G3; received basal diet till the 20th day, then the diet was coated with Difloxacin at 5 mg/kg for 10 days (Elston *et al.*, 1997) till the 30th day. One month later, the blood samples were taken from the caudal blood vessels for analysis of the blood and immune parameters.

Experiment 2: Experimental infection with *Y. ruckeri* and Difloxacin treatment (30-40 days)

It was a complementary step for the first experiment. The control G1 was subdivided equally into G1 (control negative un-injected) and G2 (control positive injected). G2, G3, and G4 were injected intraperitoneally (I/P) with 0.5 ml of a bacterial suspension of *Y. ruckeri* at 1.5×10^8 CFU /fish. Two days later, the injected groups were treated with Difloxacin 10 mg/L, one

hour bath, for 10 successive days. Along the treatment course, the clinical signs and mortalities were registered daily. Following the treatment period, the blood samples were taken and the hematological, and immunological parameters were analyzed.

Clinical signs

Either clinical internal or external abnormalities, besides, physical and behavioral signs were reported according to **Danley *et al.* (1999)**.

Blood sampling

Blood samples were drawn from the caudal vessels and directly placed in a tube containing potassium salt of EDTA and used as whole blood for hematological examination. The remaining amount of blood was placed in a plain clean tube and centrifuged at 3000 rpm for 5 minutes then the serum was separated for the biochemical and immunological assays.

Hematological examination

The erythrocytic and total leukocytic counts and the blood films were set for the differential leukocytic count with Giemsa stain according to **Blaxhall and Daisley, (1973)**. Hemoglobin concentration (Hb) was determined by the cyanomethemoglobin method (**Dorafshan *et al.*, 2008**).

Serum biochemical and immunological parameters

Serum levels of total proteins, albumin, glucose, urea, creatinine, uric acid, and transferases (ALT & AST) activities were analyzed according to **Koller, (1984)**, **Bartholomew and Delaney, (1964)**; **Young (2001)**, **Tietz (1995)**; **Bartels and Bohmer, (1972)**, and **Reitman and Frankel, (1957)** respectively. Also, lysozyme concentration was assessed by turbidimetric assay, (**Ellis, 1990**), Nitroblue tetrazolium (NBT) (**Glasser and Fiederlein, 1990**), Nitric oxide (**Green, *et al.*, 1982**). IL-2 and IL-10 were measured using ELISA kits manufactured by Biosource Inc., San Diego, California, USA.

Statistical analysis

Statistical analysis was performed on mean values using one-way analysis of variance (ANOVA) with Duncan Multiple Range Test was done to determine the differences among fish groups at the significance level of $P < 0.05$ (Duncan, 1955) by using SPSS for Windows, version 15, USA. The standard errors were estimated. Data of antimicrobial resistance was analyzed with a chi-squared test according to the SAS package (**SAS, 2005**).

RESULTS

1. Bacteriological identification

The recovered *Y. ruckeri* strains develop raised round, shiny, entire, and off-white colonies of 2–3 mm diameter. Most of the biochemical tests showed homogenous patterns except for Voges-Proskauer, gelatin, and sorbitol (**Table 1**).

2. The prevalence of *Yersinia ruckeri* infection

As shown in **Table 2**, the isolated *Y. ruckeri* strains from fish body surface and gills were reported in the five farms with a total prevalence of 7%. Except, Farm 2, the intestinal bacterial isolates recorded a 5.63% prevalence rate (**Table 3**).

Table (1): Bacteriological identification of *Y. ruckeri* isolates.

Isolates Test	Farm1	Farm 2	Farm 3	Farm 4	Farm 5
OX	-	-	-	-	-
CAT	+	+	+	+	+
MOT	+	+	+	+	+
ONPG	+	+	+	+	+
ADH	-	-	-	-	-
LDC	+	+	+	+	+
ODC	+	+	+	+	+
CIT	-	-	-	-	-
H2S	-	-	-	-	-
URE	-	-	-	-	-
TDA	-	-	-	-	-
IND	-	-	-	-	-
GLU	+	+	+	+	+
MAN	+	+	+	+	+
INO	-	-	-	-	-
RHA	-	-	-	-	-
SUC	-	-	-	-	-
MEL	-	-	-	-	-
AMY	-	-	-	-	-
ARA	-	-	-	-	-
SOR	-	+	+	-	-
G	-	+	-	-	-
VP	+	+	+	-	+
NO2	+	+	+	+	+
N2	+	+	+	+	+
O	+	+	+	+	+
F	+	+	+	+	+

OX: oxidase, CAT: catalase, β H: b haemolysis, MOT: motility, ONPG: b-galactosidase, ADH: arginine dihydrolase, LDC: lysine decarboxylase, ODC: ornithine decarboxylase, CIT: citrate, H2S: production of hydrogen sulphide, URE: urease, TDA: tryptophanedeaminase, IND: indole, VP: Voges-Proskauer, G: gelatin, GLU: glucose, MAN: mannitol, INO: inositol, SOR:sorbitol, RHA: rhamnose, SUC: sucrose, MEL: melibiose, AMY: amygdalin, ARAB: arabinose, NO2: nitrate reduction, N2: reduction to nitrite gas.

3. Antimicrobial resistance

The antimicrobial susceptibility of *Y. ruckeri* strains showed a higher resistance pattern for all antibiotics except for Difloxacin. The multi-resistant patterns of *Y.ruckeri* were shown in **Table 4**.

4. Clinical signs

The infected fish exhibited behavioral changes including surface swimming, lethargy, and decreased appetite. The common signs were bilateral exophthalmia, inflammation, erosion, darkened skin, and hemorrhages around the mouth (red mouth) also appeared. Low sustained mortalities were observed which increased over time. Internally, enlarged liver and kidney, and intestine filled with a yellowish fluid of fetid, besides, petechial hemorrhage in the liver, spleen, intestine could be noticed.

Table (2): Identification and source of selected bacterial isolates from the body surface and gills of farmed Nile tilapia (n=100 out of 5 farms).

Type of isolates	TF1	TF2	TF3	TF4	TF5	Total
<i>Yersinia ruckeri</i>	1	1	2	2	1	7

Table (3): Identification of the intestinal bacterial isolates from examined Nile tilapia (n=71).

Type of isolates	TF1	TF2	TF3	TF4	TF5	Total
<i>Yersinia ruckeri</i>	1	-	1	1	1	4

Table (4): Antimicrobial resistance of *Y. ruckeri* strains (n=56 isolates).

Bacterial strains	Antimicrobial resistance (%)						
	Farm	C	SXT	D	AMP	OT	E
F1		20	70	0	70	100	80
F2		60	40	20	70	100	90
F3		70	80	30	80	100	100
F4		40	60	30	60	80	90
F5		50	30	10	10	60	50
Chi square value		11.2**	7.5*	6.7*	0.68 ^{ns}	0 ^{ns}	4.44 ^{ns}

C: Chloramphenicol, SXT: Sulphamethoxazole/Trimethoprim, D: Difloxacin, Am: Ampicillin, OT: Oxytetracycline, E: Erythromycin. *: P value < 0.05 (significant), **: P value < 0.01 (highly significant), ns: P value > 0.05 (non-significant)

Table (5): The eight most common antimicrobial resistance profiles encountered in 81 multi-resistant *Y. ruckeri* strains from gills of Nile tilapia.

Antibiotic resistance profile	No. of <i>Y. ruckeri</i> strains
OT, SXT, AMP, E	5
OT, SXT, C	3

5. Experiment 1: Dietary supplementation by Ascorbic acid and Difloxacin

5.1. Hematological parameters

As shown in **Table 6**, Hb concentration was significantly ($P<0.05$) higher in the Ascorbic Acid-supplemented group than Difloxacin-treated group. RBCs and PCV were non significantly changed in all groups. Additionally, the supplemented groups revealed significant ($P<0.05$) leukocytosis, neutrophilia, and monocytosis in correspondence to the control.

Table (6): Some hematological parameters (mean values \pm SE) at 30th day following dietary supplementation of Nile tilapia with Ascorbic acid and Difloxacin.

Parameters	Experimental groups		
	G1 Control	G2 Ascorbic Acid	G3 Difloxacin
RBCs($10^6/\mu\text{l}$)	2.28 \pm 0.04 ^a	2.37 \pm 0.05 ^a	2.40 \pm 0.11 ^a
Hb(g/dl)	5.84 \pm 0.28 ^b	6.88 \pm 0.16 ^a	5.81 \pm 0.17 ^b
PCV(%)	21.67 \pm 0.88 ^a	21.00 \pm 0.58 ^a	20.00 \pm 0.58 ^a
MCV(fl)	95.30 \pm 4.92 ^a	88.52 \pm 4.04 ^a	83.93 \pm 6.40 ^a
MCH(pg)	25.67 \pm 1.14 ^a	28.96 \pm 0.37 ^a	24.37 \pm 1.87 ^a
MCHC(%)	27.17 \pm 2.41 ^b	32.82 \pm 1.15 ^a	29.03 \pm 0.12 ^{ab}
TLC($10^3/\mu\text{l}$)	36.21 \pm 1.23 ^b	39.68 \pm 0.49 ^a	38.67 \pm 0.29 ^{ab}
Neutrophils($10^3/\mu\text{l}$)	15.42 \pm 1.76 ^b	20.66 \pm 0.98 ^a	21.07 \pm 0.15 ^a
Lymphocytes($10^3/\mu\text{l}$)	19.27 \pm 0.51 ^a	17.28 \pm 1.01 ^{ab}	15.84 \pm 0.38 ^b
Monocytes($10^3/\mu\text{l}$)	1.18 \pm 0.10 ^b	1.47 \pm 0.07 ^a	1.55 \pm 0.08 ^a
Eosinophils($10^3/\mu\text{l}$)	0.34 \pm 0.05 ^a	0.26 \pm 0.04 ^a	0.21 \pm 0.02 ^a

RBCs: Red Blood Cells, Hb; Hemoglobin, PCV: Packed Cell Volume, MCV: Mean Corpuscular Volume, MCH: Mean Corpuscular Hemoglobin, MCHC: Mean Corpuscular Hemoglobin Concentration, TLC: Total Leucocytic Count.

5.2. Serum biochemical parameters

Similarly, the Ascorbic Acid and Difloxacin-supplemented groups showed non-significant changes in all estimated serum biochemical analytes compared to the control (**Table 7**).

Table (7): Some serum biochemical analytes (mean values \pm SE) at 30th day following dietary supplementation of Nile tilapia with Ascorbic acid and Difloxacin.

Parameters	Experimental groups		
	G1 Control	G2 Ascorbic Acid	G3 Difloxacin
AST (U/L)	7.52 \pm 0.99 ^a	8.50 \pm 0.45 ^a	9.86 \pm 0.74 ^a
ALT(U/L)	9.03 \pm 1.15 ^a	9.12 \pm 0.32 ^a	10.75 \pm 0.70 ^a
TP (g/dl)	6.00 \pm 0.09 ^a	6.11 \pm 0.03 ^a	5.39 \pm 0.44 ^a
Albumin (g/dl)	4.11 \pm 0.06 ^a	3.38 \pm 0.16 ^a	3.96 \pm 0.08 ^a
Glucose(mg/dl)	65.52 \pm 2.58 ^a	67.53 \pm 1.86 ^a	64.61 \pm 3.03 ^a
Uric acid (mg/dl)	0.95 \pm 0.06 ^a	0.94 \pm 0.03 ^a	0.94 \pm 0.04 ^a
Urea (mg/dl)	16.72 \pm 0.59 ^a	18.18 \pm 0.95 ^a	18.58 \pm 1.18 ^a
Creatinine (mg/dl)	43.92 \pm 2.80 ^a	41.71 \pm 2.18 ^a	43.77 \pm 2.27 ^a

AST: Aspartate aminotransferase, ALT: Alanine aminotransferase, TP: Total Protein.

5.3.The immunological assays

Significant ($P<0.05$) elevations were showed in Lysozyme, NO, Phagocytic activity %, IL2, IL10, and SOD concentration in the Ascorbic Acid and Difloxacin- supplemented groups compared to the control. While NBT was non significantly increased in the supplemented groups (Table 8).

Table (8): Some immunological assays (mean values \pm SE) at 30th following dietary supplementation of Nile tilapia with Ascorbic acid and Difloxacin.

Parameters	Experimental groups		
	G1 Control	G2 Ascorbic Acid	G3 Difloxacin
NBT(OD)	0.96 \pm 0.05 ^a	0.99 \pm 0.03 ^a	0.97 \pm 0.03 ^a
Lysozyme(U/L)	82.12 \pm 3.62 ^b	109.75 \pm 6.05 ^a	103.77 \pm 4.63 ^a
NO(μ g/dl)	25.28 \pm 1.58 ^b	41.67 \pm 1.31 ^a	37.13 \pm 2.36 ^a
SOD(U/ml)	249.08 \pm 2.95 ^b	270.41 \pm 4.29 ^a	261.05 \pm 3.55 ^a
Phagocytic(%)	52.38 \pm 1.87 ^b	83.63 \pm 2.18 ^a	75.93 \pm 3.00 ^a
IL-2(Pg/ml)	4.19 \pm 0.22 ^c	5.18 \pm 0.24 ^b	7.72 \pm 0.23 ^a
IL-10(Pg/ml)	3.11 \pm 0.13 ^c	4.04 \pm 0.22 ^b	6.38 \pm 0.21 ^a
Mortality rate	3.33 \pm 1.67 ^a	1.67 \pm 1.67 ^a	3.33 \pm 3.33 ^a

NBT: Nitroblue tetrazolium, NO: Nitric Oxide, SOD: Superoxide Dismutase, IL: Interleukin.

6. Experiment 2: Experimental infection with *Yersinia ruckeri* and Difloxacin treatment**6.1. Hematological parameters**

On the 40th day, the *Yersinia*-infected fish revealed a significant decrease in RBCs, Hb content, and PCV compared to the control and treated fish (**Table 9**).

6.2. Serum biochemical parameters

A significant ($P<0.05$) increase in AST activity in *Yersinia*-infected fish compared to the control. Meanwhile, the treated groups showed a numerical decline in AST activity compared to the infected group. Furthermore, the infected group manifested a significant ($P<0.05$) increase in ALT activity and urea level in comparison to the control and treated groups. Significant ($P<0.05$) hypoproteinemia and hypoalbuminemia were detected in the infected group in comparison to the control and treated groups. As for creatinine, uric acid, and glucose levels, there were non-significant variations between all groups, except for a numerical increase in creatinine level in *Yersinia* infected group (**Table 10**).

Table (9): Some hematological parameters (mean values \pm SE) at 40th day in the healthy control and *Y. ruckeri*-infected fish supplemented with Ascorbic acid and Difloxacin.

Parameters	Experimental groups			
	G1	G2	G3	G4
RBCs($10^6/\mu\text{l}$)	2.52 \pm 0.18 ^a	1.21 \pm 0.06 ^b	2.43 \pm 0.08 ^a	2.41 \pm 0.06 ^a
Hb(g/dl)	6.51 \pm 0.16 ^a	5.18 \pm 0.09 ^b	7.26 \pm 0.77 ^a	6.82 \pm 0.06 ^a
PCV(%)	21.33 \pm 0.67 ^a	16.00 \pm 0.58 ^b	21.67 \pm 0.88 ^a	20.67 \pm 1.45 ^a
MCV(fl)	84.75 \pm 2.25 ^b	133.08 \pm 6.13 ^a	89.17 \pm 3.97 ^b	86.02 \pm 7.49 ^b
MCH(pg)	25.89 \pm 0.78 ^b	43.06 \pm 1.50 ^a	29.71 \pm 2.13 ^b	28.31 \pm 0.46 ^b
MCHC(%)	30.64 \pm 1.73 ^a	32.41 \pm 0.88 ^a	33.50 \pm 3.05 ^a	33.35 \pm 2.57 ^a
TLC($10^3/\mu\text{l}$)	37.24 \pm 0.55 ^{ab}	30.87 \pm 2.14 ^b	35.15 \pm 2.87 ^{ab}	39.97 \pm 2.70 ^a
Neutrophils($10^3/\mu\text{l}$)	14.78 \pm 0.24 ^{ab}	9.40 \pm 0.45 ^b	19.62 \pm 2.04 ^a	19.49 \pm 2.98 ^a
Lymphocytes($10^3/\mu\text{l}$)	21.00 \pm 0.27 ^a	19.96 \pm 1.62 ^a	13.92 \pm 0.95 ^b	18.60 \pm 0.36 ^a
Monocytes($10^3/\mu\text{l}$)	1.05 \pm 0.08 ^b	1.15 \pm 0.08 ^b	1.32 \pm 0.08 ^{ab}	1.57 \pm 0.19 ^a
Eosinophils($10^3/\mu\text{l}$)	0.40 \pm 0.04 ^a	0.36 \pm 0.01 ^{ab}	0.28 \pm 0.02 ^b	0.30 \pm 0.01 ^b

RBCs: Red Blood Cells, Hb: Hemoglobin, PCV: Packed Cell Volume, MCV: Mean Corpuscular Volume, MCH: Mean Corpuscular Hemoglobin, MCHC: Mean Corpuscular Hemoglobin Concentration, TLC: Total Leucocytic Count. **G1:** Control un- challenged group, **G2:** Control challenged group, **G3:** challenged Ascorbic Acid-treated group, and **G4:** challenged Difloxacin treated group.

Table (10): Some serum biochemical analytes (mean values \pm SE) at 40th day in healthy control and *Y. ruckeri*-infected fish supplemented with Ascorbic acid and Difloxacin.

Parameters	Experimental groups			
	G1	G2	G3	G4
AST (U/L)	9.68 \pm 0.62 ^b	15.57 \pm 1.91 ^a	10.90 \pm 1.23 ^{ab}	13.53 \pm 2.31 ^{ab}
ALT(U/L)	11.98 \pm 1.17 ^b	22.38 \pm 1.55 ^a	11.59 \pm 0.75 ^b	11.18 \pm 1.61 ^b
TP (g/dl)	5.83 \pm 0.17 ^a	3.50 \pm 0.32 ^b	5.22 \pm 0.80 ^a	5.32 \pm 0.63 ^a
Albumin (g/dl)	5.33 \pm 0.19 ^a	2.18 \pm 0.22 ^b	4.87 \pm 0.11 ^a	4.98 \pm 0.51 ^a
Glucose (mg/dl)	50.82 \pm 2.67 ^a	64.87 \pm 3.35 ^a	55.06 \pm 6.05 ^a	52.51 \pm 4.49 ^a
Uric acid (mg/dl)	1.26 \pm 0.19 ^{ab}	1.41 \pm 0.27 ^a	0.79 \pm 0.05 ^b	0.94 \pm 0.05 ^{ab}
Urea (mg/dl)	15.09 \pm 0.28 ^b	22.80 \pm 1.77 ^a	17.52 \pm 0.84 ^b	18.19 \pm 0.50 ^b
Creatinine (mg/dl)	37.47 \pm 0.26 ^a	48.16 \pm 4.80 ^a	39.04 \pm 1.67 ^a	45.38 \pm 3.90 ^a

AST: Aspartate aminotransferase, ALT: Alanine aminotransferase, TP: Total Protein. **G1**: Control un-challenged group, **G2**: Control challenged group, **G3**: challenged Ascorbic Acid-treated group, and **G4**: challenged Difloxacin treated group.

Table (11): Some immunological assays (mean values \pm SE) at 40th day in healthy control and *Y. ruckeri*-infected fish supplemented with Ascorbic acid and Difloxacin.

Parameters	Experimental groups			
	G1	G2	G3	G4
NBT(OD)	0.92 \pm 0.02 ^a	0.78 \pm 0.02 ^b	0.97 \pm 0.03 ^a	0.90 \pm 0.01 ^a
Lysozyme(U/L)	82.09 \pm 1.66 ^a	10.28 \pm 0.65 ^c	53.14 \pm 2.11 ^b	52.21 \pm 1.60 ^b
NO(μ g/dl)	21.74 \pm 1.36 ^a	2.05 \pm 0.38 ^c	15.68 \pm 0.90 ^b	13.02 \pm 1.48 ^b
SOD(U/ml)	246.43 \pm 3.81 ^b	218.52 \pm 7.46 ^c	266.04 \pm 3.44 ^a	255.88 \pm 3.84 ^{ab}
Phagocytic(%)	41.85 \pm 1.40 ^b	21.49 \pm 1.29 ^c	72.60 \pm 1.85 ^a	73.96 \pm 2.82 ^a
IL-2(Pg/ml)	5.00 \pm 0.45 ^b	7.14 \pm 0.21 ^a	4.82 \pm 0.20 ^b	3.87 \pm 0.19 ^c
IL-10(Pg/ml)	4.53 \pm 0.37 ^b	5.79 \pm 0.15 ^a	3.58 \pm 0.30 ^c	3.25 \pm 0.14 ^c
Mortality rate	1.67 \pm 1.67 ^c	61.67 \pm 6.01 ^a	36.67 \pm 3.33 ^b	38.33 \pm 3.33 ^b

NBT; Nitroblue tetrazolium, NO; Nitric Oxide, SOD; Superoxide Dismutase, IL; Interleukin. **G1**: Control un-challenged group, **G2**: Control challenged group, **G3**: challenged Ascorbic Acid-treated group, and **G4**: challenged Difloxacin treated group.

6.3. The immunological assays

In the *Yersinia*-infected group, NBT, Phagocytic %, lysozyme activity, NO level and SOD activity were significantly ($P < 0.05$) decreased and IL2 and IL10 were significantly ($P < 0.05$) increased in comparison to the control and treated groups (**Table 11**).

DISCUSSION

Throughout this study, some of the epidemiological characteristics of *Yersinia ruckeri* infection in Nile tilapia *Oreochromis niloticus* such as occurrence, outbreaks, and prevalence in organs were known. The behavior of the disease was also defined experimentally, in terms of its clinical signs, postmortem, and mortality rate. More importantly, is the evaluation of the hematological, biochemical, and immunological activities, during the infection, and post-treatment.

Yersinia, a genus of the Enterobacteriaceae family, comprises several species, which cause diseases in humans, animals, and fish. The infected fish exhibited some behavioral, clinical, and postmortem changes which were most similar to the previously described by **Danley et al. (1999)**.

The bacteriological identification of bacterial isolates manifested Gram-negative bacilli, oxidase-negative, catalase-positive glucose and mannitol fermenter, nitrate-reductive, secrete lysine, b-galactosidase, and ornithine decarboxylases but neither indole nor hydrogen sulfide. Similar biochemical patterns were identified by **Furones et al. (1993)** and **Elmeer et al. (1997)**. Also, the identified strains from different farms were largely homogenous, with some biochemical variation that might be related to the genetic diversity within *Y. ruckeri* strains (**Bastardo et al., 2011**).

The spatial occurrence of *Y. ruckeri* infection showed significant prevalences in the different farms. This suggests the epizootic of the infection and the high adaptation of exploding to be widely distributed, moreover, this assures the evolutionary relationships among diverse bacterial species. Thus *Y. ruckeri* has a wide range of locality and host that could explain their ability to persist longer in the aquatic environment (**Romalde et al., 1994**), and transmission through carriers to the healthy fish at an optimum temperature under any stressors (**Hunter et al., 1980**).

By screening the susceptibility of *Y. ruckeri* isolates, they were showed resistance patterns to OT, SXT, AMP, and E. Similar resistance was interpreted by **Rodgers, (2001)** in oxolinic acid, oxytetracycline, and potentiated sulphonamide. He has attributed it to *Y. ruckeri* β -lactamase gene which is not likely to be expressed at high levels (**Stock et al., 2002; Mammeri et al., 2006**). Oppositly, **Klein et al. (1996)** did not found resistance patterns among *Y. ruckeri* isolates, except for a lower sensitivity to OXA. Thus, the pathogenic strains evolve different antimicrobial resistance due to misuse of antimicrobial agents.

Many factors could be contributed to pathogenicity and the appearance of external, internal signs, and mortalities. They could be related to the generalized septicemia and the widespread of the pathogen in all organs, including the brain (**Ohtani et al., 2014**). **Méndez and Guijarro, (2013)** revealed that some promoters such as *yhlBA*, *cdsAB*, *yctCBA*, and *yyp1* were expressed and led to hemolysin production, cysteine and citrate uptake, and the encoding of serralyisin metalloprotease, respectively. Other factors have been linked to the pathogenicity such as a serralyisin metalloprotease (metalloendopeptidase), termed Yrp1, that hydrolyses actin, fibrinogen, gelatin, laminin, and myson (but not type II and type IV collagen) (**Fernandez-Diaz et al., 2003**).

The hematological parameters of yersinia-infected fish recorded macrocytic normochromic anemia. Our result came in agreement with **Altun and Diler, (1999)** who reported macrocytic normochromic anemia in rainbow trout experimentally infected with *Y. ruckeri* on day 13 and 15. Macrocytic anemia takes place due to the failure of DNA division of the erythrocytic cell (**Hoffman *et al.*, 2009**).

Vitamin C is a necessary immunostimulant facing bacteria and virus-infected fish and is responsible for the proliferation and phagocytosis of the immune cells (**Ergönül *et al.*, 2012**). **Tewary and Patra, (2008)** have concluded that high doses of vitamin C (500 to 2000 mg/kg) in rainbow trout stimulated hematopoiesis, raised the immune response, and repaired tissue damage against most viral and bacterial pathogens including *Y. ruckeri*.

At the same time, an increase mainly in serum activities of ALT, AST, and level of urea was reported along with hypoproteinemia and hypoalbuminemia in the infected non-treated group. These marked changes could be attributed to the marked degenerative changes that have been noticed in the infected tissues of the liver and kidneys (**Salah *et al.*, 2012**). It has been previously discussed that the increased serum liver transaminases (ALT and AST) may elucidate hepatocellular injury and inflammatory reaction leading to the emission of these enzymes into the blood (**Fuchs *et al.*, 1986**). Furthermore, serum ALT and AST activities are counted as sensitive signals to estimate hepatic and myocardial deterioration (**Raa, 1994**). At the same pace, the obtained results of serum hypoproteinemia and hypoalbuminemia confirmed the occurrence of liver disorder due to stress conditions and bacterial infection (**Ellis, 1981**).

Moreover, some immunological assays such as NBT, Phagocytic %, NO, Lysozyme, and SOD activities showed declination in the infected group. The potency of the phagocytic cells neutrophils, monocytes, and macrophages is associated with activation of the innate immune response (**Ayub *et al.*, 2003**). Neutrophils are an essential partner of the host defense and valuable for the assessment of the fish's health status. The macrophages are considered a focal part of the cellular non-specific defense. An inflammatory reaction has been shown in the body cavity in the *Y. ruckeri* infected rainbow trout (**Afonso *et al.*, 1998**), due to the flow of large numbers of neutrophils, caught from the blood and hemopoietic tissue. Neutrophils actively share in the nonspecific immunity of the fish by producing reactive oxygen species or radicals for killing bacteria, then Nitro-blue tetrazolium (NBT) reacts with these oxidative radicals producing a dark blue stain (**Anderson *et al.*, 1992; Barman *et al.*, 2013**). Therefore, the phagocytic activity of neutrophils can be positively correlated to the NBT activity.

Another defense element, Superoxide dismutase enzyme (SOD) which is detected in erythrocytes and the liver and prevent oxidative fatigue of the cells. High SOD activity protects the cells against oxidative stress caused by bacterial infections. Where, it is concerned with defensive mechanisms against oxidative stress as that provoked by intracellular pathogens like *Y. ruckeri* (**Guijarro *et al.*, 2018**).

Nitric oxide (NO) is bactericidal reactive oxygen that is produced by phagocytes (macrophages, monocytes, and neutrophils) following microbial invasion (**Campos-Perez *et al.*, 2000**). NO was found to cause DNA damage of the pathogen (Juedes and Wogan, 1996) and adjust the immune response toward the pathogen.

Lysozyme is an index of the innate immunity of the fish and one of the defense mechanisms of the body toward microbial invasion, due to its anti bactericidal impact (**Jollès and Jollès, 1984**) against different bacterial species (**Saurabh and Sahoo, 2008**). It can break the chemical bond between the N-acetylglucosamine and N-acetylmuramic acid of the peptidoglycan in bacterial cell walls, and mainly secreted by macrophages and neutrophils (**Nathan, 1987**). **Jaafar et al. (2013)** stated that experimentally infected fish with *Y. ruckeri* exhibited a reduction in the lysozyme activity after 45 days and this reduction may be attributed to exhaustion of the plasma lysozyme in fish blood facing the bacterial invasion. In a challenge test with vibriosis, a negative correlation between serum lysozyme activity and survival has been observed in Atlantic salmon (**Roed et al., 1993**).

On the other hand, ascorbic acid and antibiotic supplementation to the infected groups resulted in enhanced activity of lysozymes in comparison to the infected non-treated group. Increased lysozyme activity can be attributed to the dietary supplements for fish, through the synergistic effect on the immune system due to the increase in the phagocyte number elevates lysozyme synthesis by cells (**Kumari and Sahoo, 2006**).

Furthermore, IL2 and IL10 showed an elevation in their levels in the infected non-treated group. The inflammatory response in *Y. ruckeri* infected rainbow trout reported raised levels of inflammatory cytokines such as interleukin (IL)-1 family members, IL-6, IL-8, IL-10, and IFN- γ (**Raida and Buchmann, 2008; Wang et al., 2009; Kumar et al., 2015**). Interleukin IL-2 is an immunomodulatory element that helps proliferation, differentiation, and activation of T cells (**Sogo et al., 2009**). It is synthesized mainly by Th1 cells and is known as the T-cell growth factor (TCGF) (**Smith, 1988**). Also, the expression of Interleukin IL-10 can be increased by a bacterial infection, LPS stimulation, and immunostimulants bath supplementation (**Zhang et al., 2009**). IL-10 depresses iNOS and thus downregulates NO production, resulting in disease promotion (**Huang et al., 1998**). **Raida and Buchmann, (2008)** found that *Y. ruckeri* O1 amount in the spleen was correlated to the expression of cytokines genes 1 β , IL-8, and IL-10 in rainbow trout. Moreover, the infected fish showed a significant rise in the transcript levels of chemokines, central cytokines, and cytokine receptors (IL-1 β , IL-6, IL-8, IL-10, TNF- α , IL-receptor II) (**Raida et al., 2011**).

CONCLUSION

Thus it could be concluded that the pathogenesis of Yersiniosis in the cultured *Oreochromis niloticus* has been investigated in terms of clinical signs, pathogenicity patterns, and disturbances of hematological, and immunological responses. Also, Ascorbic Acid supplementation and Difloxacin dosing have proven their efficiency against *Yersinia ruckeri* infection indicated in improving health status and enhancement of biochemical and immune responses. Therefore ascorbic acid should be included within the preventive programs.

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