



A Study of the Effect of Selenium Nanoparticles on *Aeromonas* Resistance in Nile Tilapia Fish

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Abstract

Aeromonas hydrophila is a significant pathogen affecting warm-water fish, particularly Nile tilapia, leading to various bacterial illnesses. This study investigates the limitations of antibiotics in treating *A. hydrophila* infections and explores selenium nanoparticles (Se-NPs) as a potential alternative treatment. The research was conducted in Sharkia governorate, Egypt, where a high prevalence (92%) of *Aeromonas* spp. was found among isolated strains, all exhibiting multidrug resistance to 11 out of 13 antibiotics tested. In a 35-day feeding trial involving 120 Nile tilapia, fish were divided into four groups: a control group, a group receiving a basal diet supplemented with Se-NPs, a positive control group infected with *A. hydrophila*, and a group receiving Se-NPs while also infected. Results indicated that *A. hydrophila* infection led to significant alterations in hematological and biochemical profiles, including increased levels of liver enzymes (ALT, AST, ALK, GGT), urea, creatinine, and uric acid, alongside decreased total protein, albumin, globulin, and antioxidant enzyme levels. However, dietary supplementation with Se-NPs significantly improved these parameters, enhancing antioxidant activity and immune response in Nile tilapia. Furthermore, Se-NPs mitigated the pathological effects caused by *A. hydrophila* infection, suggesting their potential as an effective treatment strategy for managing bacterial diseases in aquaculture.

Keywords: *Oreochromis niloticus*, Selenium nanoparticles, *Aeromonas hydrophila*, hematology. Antioxidants, Phagocytic percent.

Introduction

One of the animal food production sectors with the greatest rate of growth in the world is aquaculture, which significantly boosts both economic growth and global food security [1]. The Nile tilapia, (*Oreochromis niloticus*), is a common species of freshwater fish because of its rapid growth, ability to adapt to a variety of environmental circumstances, and high demand from consumers [2]. However, intense aquaculture operations frequently expose fish to stress and opportunistic bacterial infections, resulting in substantial economic losses. *Aeromonas*

species, notably *A. hydrophila*, are recognized as the primary.

Causal agents of motile *Aeromonas* septicemia (MAS), characterized by hemorrhages, fin rot, ulcers, and high mortality rates [3]. Bacterial infections in fish are frequently associated with oxidative stress, which is caused by a surplus of reactive oxygen species (ROS) that overwhelm the antioxidant defense systems. Oxidative damage can harm key organs, including the liver and kidneys, affect protein metabolism, and weaken immunological responses [4]. Thus, nutritional measures aimed at increasing antioxidant capacity and immune defense are

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regarded as useful complementary approaches to disease control in aquaculture. In this setting, there is a critical need for environmentally benign, effective antibiotic alternatives that can improve disease resistance and immune response in fish. An interesting solution is to use nanotechnology, specifically nano-selenium (Se-NPs) as a food supplement in aquaculture.

Selenium (Se) is an essential trace element that aids the antioxidant defense system. It is largely a component of selenoenzymes such as glutathione peroxidase (GPx), which guards against oxidative cell damage [5]. Selenium has been shown to enhance immune responses and disease resistance in fish, in addition to its antioxidant properties [6]. However, the bioavailability of traditional selenium sources is frequently limited, and excessive supplementation may be harmful [7].

Nano-selenium, a unique form of selenium with particle sizes in the nanometer range, has received interest in aquaculture nutrition due to its better bioavailability, lower toxicity, and robust biological activity compared to inorganic and organic selenium forms [8]. Recent studies have revealed that dietary nano-Se can enhance growth, boost antioxidant status, and protect against bacterial and environmental stresses in fish [9, 10].

However, there has been little investigation into its therapeutic efficacy against *Aeromonas* infections in Nile tilapia, notably in terms of liver and kidney function, protein metabolism, and antioxidant responses. Therefore, the current work was conducted to study the protective effects of dietary Se-NPs on liver and kidney biomarkers, serum protein profile, and antioxidant status in Nile tilapia experimentally infected with *Aeromonas* spp. The findings aim to shed light on the potential application of nano-selenium as a feed additive in aquaculture to enhance health and resistance to diseases.

Material and Methods

Fish

An earthen fish farm at the Central Laboratory for Aquaculture Research in Abassa, Sharkia province, Egypt, donated 120 healthy male Nile tilapia weighing 30 ± 1.8 g to the Animal Health Research Institute's Zagazig branch. The fish were transported in plastic bags containing water. Fish were acclimatized for two weeks in glass tanks ($100 \times 40 \times 40$ cm) with 100 liters of dechlorinated tap water. The tanks were aerated using an electric air pump, and heaters were utilized to keep the natural photoperiod temperature of 26°C. The water was maintained at 6.5 ± 0.5 , 7.1 ± 0.8 , 0.003, 0.31, and 0.42 mg L⁻¹ for dissolved oxygen, pH, ammonia, nitrate, and nitrite levels. A basal diet containing 30% protein was administered twice daily at a rate of 4%

of body weight for the duration of the two-week adaptation period.

Selenium nanoparticles

Mint leaf extract was used as a stabilizing and reducing agent in an environmentally friendly reduction procedure to create green selenium nanoparticles. After being thoroughly cleaned with deionized water and chopped into small pieces, fresh mint leaves were boiled for 20 minutes at 80 degrees Celsius in 100 milliliters of distilled water. Whatman filter paper was used to filter the extract, which was then stored for later use at 4°C. Mint extract was added dropwise to a sodium selenite (Na₂SeO₃) solution while being continuously stirred at room temperature to create the selenium nanoparticles. As soon as selenium nanoparticles were formed, the reaction mixture's color changed from colorless to brick red. For four hours, the mixture was constantly agitated to guarantee full reduction and stabilization. To obtain selenium nanoparticle powder, the resultant nanoparticles were collected by centrifugation at 10,000 rpm for 15 minutes, repeatedly cleaned with distilled water and ethanol to get rid of any materials that hadn't reacted, and then dried in a vacuum oven at 60°C for 12 hours [11-14]. UV-visible spectra and XRD patterns were used to identify the produced Se-NPs and determine their size, shape, and morphology [13, 15, 16 & 17].

Experimental Design

Four groups of thirty fish each were created from 120 freshwater Nile tilapia. The fish were randomly divided into two replicates of four groups, 15 fish each. Each group received one of the treatments indicated below:
Group 1 (G1): Fish were fed a baseline diet (30% crude protein) for 35 days and served as the control.
Group 2 (G2): Fish were fed the base diet supplemented with Nano-selenium 0.3 mg/kg according to [18] from day one to the completion of the experiment (35 days).
Group 3 (G3): Fish were fed a basal diet from day one to the completion of the trial (35 days) and artificially infected with *Aeromonas hydrophila* on day 21 of the experiment. Each fish received a 0.5 ml intraperitoneal injection of 1×10^7 CFU and served as a positive control, following [19].
Group 4 (G4): fish were fed a basal diet supplemented with Nano-selenium 0.3 mg/kg from day one to the completion of the experiment (35 days) and artificially infected with *Aeromonas hydrophila* on day 21 of the experiment.

For 35 days, each group's fish were given 3% of their body weight twice a day. During the acclimation stage, half of the water was changed daily to avoid the buildup of feces. The test included an analysis of the water quality measures. Collecting samples after 35 days from all groups, 14 days after infection in groups 3 and 4.

Bacterial Isolation

Seemingly healthy and diseased Nile tilapia were collected from the market place in Sharkia Governorate (muscles, fins, intestine, gills, and skin) during the summer period (2024). Fish specimens were promptly and aseptically conveyed to the lab of the Microbiology Department, Animal Health Research institute. All fish exhibited indications of hemorrhages around the operculum region (circle) and ulceration on the tail fin.

Conventional identification

The specimens were cultivated in nutrient broth (Oxoid), and kept aerobically for 18–24 hours at 37°C. After being streaked across *Aeromonas* agar base medium (Hi-medium, India). A single colony from the agar was used to make a smear stained with Gram stain and then morphologically observed under a microscope [21].

Re-isolation of A. hydrophila was made from all groups

Molecular identification of *A. hydrophila*

The QIAamp DNA Mini kit (Qiagen, Germany, GmbH) was used to extract DNA from samples, with some adjustments made based on the manufacturer's instructions. Metabion (Germany) provided the primers (Table 1).

PCR amplification

A 25 µl reaction comprising 12.5 µl of Emerald Amp Max PCR Master Mix (Takara, Japan), 1 µl of each primer at a concentration of 20 pmol, 5.5 µl of water, and 5 µl of DNA template was used to use primers. An Applied Biosystem 2720 heat cycler was used to carry out the process.

Analysis of the PCR Products

A 1.5% agarose gel was used for electrophoresis of the PCR product. A gel documentation system (Alpha Innotech, Biometra) took pictures of that, and a computer was used to analyze the data.

Sensitivity assay

Examination of Antimicrobial activity of A. hydrophila for Se NPs

The disc diffusion approach utilized. Sterile discs (0.6mm diameter) of Whatman No. 2 filter paper were loaded with 100 µL/disc of Se NPs at a concentration of 10000 µg/ml and positioned on Muller Hinton agar plates inoculated with bacterial cultures, generated inoculum equivalent to 0.5 McFarland (1.5 x 10⁸ CFU). Petri plates incubated at 37 degrees Celsius for 24 hours, and the inhibitory zone was properly quantified [23].

Antimicrobial susceptibility testing of Aeromonas spp for antimicrobial agents

Using the disc diffusion method and the CASFM recommendations, all of the isolates were evaluated for resistance to 13 antimicrobial drugs (Table 2). [24] regarding its susceptibility to antimicrobials [(Oxoid, UK). Some colonies were moved to a tube containing 5 ml of sterile physiological saline and matched with a 0.5 McFarland standard tube (1.5 x 10⁸ CFU). Antimicrobial discs were put on the Muller Hinton agar after the bacterial culture had been scattered onto the plates surface. Afterward, incubation at 37°C for 24 hours. The outcomes were interpreted by CLSI [25].

Sampling

After a 24-hour fast, five fish from each group were chosen at the end of the experiment and given a 5-minute anesthesia with buffered tricaine-methane sulfonate (E10521-10G, Sigma-Aldrich) at a dosage of 100 mg/L [26]. Blood samples were obtained from caudal blood vessels and placed in a 1-mL sterile syringe with 0.2 mL of EDTA for hematological parameter evaluation, heparinized tubes for phagocytic test, and dry, clean centrifuge tubes for serum separation. After 15 minutes at ambient temperature, the tubes were centrifuged for 10 minutes at 3000 rpm.

Hematological studies:

A Neubauer's hemocytometer was used to manually estimate the total erythrocyte count (RBC × 10⁶/µl) and total leukocyte count (WBC × 10³/µl), with Hayem solution serving as the diluent [27]. The cyanmethemoglobin technique, [28] was used to compute hemoglobin (Hb), and the hematocrit % was determined using a microhematocrit centrifuge [29]. Mean corpuscular volume (MCV), mean corpuscular hemoglobin (MCH), and mean corpuscular hemoglobin concentration (MCHC) were calculated [30].

Biochemical studies

Commercial kits and established protocols were used to determine the serum biochemical parameters. Using spectrophotometric kits (Biomed Diagnostic, Egypt), the activities of aspartate aminotransferase (AST) and alanine aminotransferase (ALT) were measured [31, 32]. The activities of alkaline phosphatase (ALP) and gamma-glutamyl transferase (GGT) were examined [33]. The concentrations of albumin (ALB) and total protein (TP) were determined [34], and the difference between the ALB and TP values was used to compute globulin (GLOB) [35].

Renal function markers were analyzed following standard protocols. Serum creatinine concentration was determined according to the colorimetric method [36]. Urea levels were measured [37], while uric acid concentration was estimated according to the enzymatic procedure [38].

Antioxidant enzyme activities were evaluated to monitor oxidative stress responses. Superoxide dismutase (SOD) activity was measured using both an ELISA kit (Catalog No. CSB-E08555r, Cusabio Biotech Co., China) and the NADH oxidation method [39]. Glutathione peroxidase (GPx) activity was determined [40], and catalase (CAT) activity was assayed [41].

Phagocytic % and phagocytic index

By determining the percentage of macrophages in a random sample of 300 macrophages that house intracellular yeast cells and expressing the result as a percentage of phagocytic activity, phagocytosis was evaluated. The phagocytosis assay will be computed [42] using the following equations:

Phagocytic activity is calculated as yeast-containing macrophages divided by the total number of macrophages $\times 100$.

Phagocytic index equals the number of yeast cells phagocytized divided by the number of phagocytic cells.

Statistical analysis

The One-Way Analysis of Variance (ANOVA) test was performed to statistically analyse the data. SPSS 14.0 (2006) was used to perform the Duncan test on data reported as mean $\pm$ standard error (SE). Statistical significance was defined as $p < 0.05$.

Results

Phenotypic and genotypic identification of the bacterial isolate

Aeromonas spp detected with an incidence rate of 92% of the bacteriologically tested samples. *Aeromonas* spp showed round, smooth colonies with 2-3 mm size and opaque green with a dark center on *Aeromonas* medium base (Fig. 1). Under the microscope, it revealed gram-negative, straight rods bacteria with rounded ends (bacilli to coccobacilli shape), motile by a single polar flagellum. Confirmation by molecular technique observed the existence of *A. hydrophila* 16S rRNA gene utilizing a specific primer for species identification yielding amplicon size 685 bp (Fig. 2).

Re-isolation of the inoculated bacterial isolate was obtained from the experimentally infected fish (G3,G4) and the culture and molecular examination of the re-isolated *A. hydrophila* revealed the same phenotypic and genotypic characteristics of the inoculated one, while re-isolation of *A. hydrophila* from G1,G2 show negative results.

Antimicrobial susceptibility findings

A surprisingly elevated proportion of resistance to diverse antimicrobial agents and development of MDR among the *A. hydrophila* strains was observed (Fig. 3 a). All isolated strains demonstrated 100%

resistance to all used antibiotics with the exception of Gentamicin, they showed 66.6% resistance, and 33.3% were sensitive and for Neomycin, 75% of isolates were resistant, whereas 25% were intermediate (Table 3).

On the other hand, findings of disc diffusion technique used to determine the zone of inhibition after exposure of *A. hydrophila* to Se NPs showed inhibition zones ranged from 11mm: 28mm diameter (Fig. 3 b).

Hematological results

The results, shown in Table (4), revealed a significant rise in total erythrocytic count, hemoglobin content, and packed cell volume percent in non-challenged groups supplemented with Se-NPs (group 2) compared to the control group. However, after *A. hydrophila* infection, there was a substantial decrease in RBC count, hemoglobin concentration, and hematocrit %, as well as a significant increase in MCV in group 3, compared to the control. Group 4, challenged with *A. hydrophila* and supplemented with Se-NPs, showed an improvement in all hematological parameters compared to the challenged non-treated group (G 3).

Moreover, the Se-NPs supplemented group (G2) showed significant increase in total leukocytic, neutrophilic, lymphocytic, and monocytic counts compared to the control group as recorded in Table (5). Infected group with *A. hydrophila* revealed changes in leukogram represented by a significant increase in total leukocytes, neutrophils and monocyte count with a significant decrease in lymphocytic count compared with groups 1 and 2. Eosinophils and basophils counts revealed no significant difference between all experimental groups

Biochemical results

The results in Table (6) demonstrated that the Se-NPs supplemented group (G2) had no significant differences in liver or kidney functions when compared to the control. *Aeromonas*-infected group (G3) demonstrated significant increases in ALT, AST, ALK, GGT activity, urea, creatinine, and uric acid levels, linked with a significant drop in total protein, albumin, and globulin levels compared to either the control group or the Se-NPs-treated group.

Oxidative stress responses

As shown in Table (7), SOD, CAT, and GPx activity were highest in G2 and lowest in G3 (infected control) when compared to control, although G4 exhibited considerably enhanced SOD, CAT, and GPx activity relative to G3 but did not reach the levels of G1 or G2.

Phagocytic Activity

Phagocytic percentage and phagocytic activity increased significantly in all experimental groups compared to the control, with the greatest levels in groups 3 and 4, as shown in Table (7).

Discussion

Fish is one of the most fundamental foods because of its great nutritional value, delicious flavor, and ease of digestion. However, it acts as a source of many harmful bacteria, particularly *Aeromonas* spp., which may pose public health risks. Fish bacterial contamination is blamed on a variety of sources, including water, soil, and handlers. Poor handling during catching, transporting, and freezing can lead to fish contamination from the environment. These bacteria are linked to a variety of human ailments, making aquaculture products a possible danger factor for customers [43].

The present study explored the prevalence of *Aeromonas* spp. in *Tilapia nilotica* fish in Egypt. High proportion (92%) of the examined fish contaminated with *Aeromonas* spp, perhaps during storage, transit or merchandising, these findings concur in some measure with studies conducted in Turkey [44] and Malaysia [45], in which *Aeromonas* spp. was isolated in a high percentage of 82.8% and 69% of fish samples respectively. The 16S rRNA gene provides an effective molecular identification and a quick technique to assess the identity of *A. hydrophila* [46].

The current study has proven the existence of MDR *Aeromonas* spp. Also, the high MDR indices of *Aeromonas* spp were detected in Nile tilapia broodstock [47].

A. hydrophila was highly resistant to the thirteen antimicrobial agents used in the present investigation. The isolates were 100% resistant to all used antibiotics (DO/30), (SXT/1.25 /23.75), (CIP₅), (C/30), (E/15), (S/10), (K/30), (AX/25), (NA/30), (CT/10 (FEP/30) , except for Gentamicin(CN/10), they showed 66.6% resistance and 33.3% were sensitive and Neomycin (N/30) 75% of isolates were resistant while 25% were intermediate, similar results were obtained by another study in Egypt. As *A. hydrophila* isolates were resistant to erythromycin and streptomycin in percentages of (100% each) but it Conflicts with our findings as it was sensitive to ampicillin (80%) and gentamicin (60%).

In the current study, results of the disc diffusion technique applied after exposure of *A. hydrophila* to Green selenium nanoparticles synthesized from mint leaf extract, Se NPs showed inhibition zones ranged from 11mm : 28mm diameter. Similarly, the Se NPs synthesized from *Blumea axillaris* plant inhibited *Aeromonas* species with an inhibition zone size of 1.8 cm (stem) and 2.6 cm(root). Also, it inhibited *A. hydrophila* (moderate), with inhibition zone size of 2.8 cm (stem) and 3.0 cm (root), whereas it was 1.0

cm (stem) and 1.5 cm(root) against *A. hydrophila* (virulent) y [48].

Se-NPs exhibited efficacy as antimicrobial agent at a concentration of 10000µg/ ml via disk diffusion technique, against *A. hydrophila* in our study. Also, antibacterial activity of SeNPs was confirmed with different concentrations (5000, 4000, 3000, 2000, 1000, 900, 800, 700, 600, 500, 400) µg/mL which inhibited the growth of many pathogenic Gram-negative and Gram-positive bacteria (*S. typhi*, *E. coli*, *P. aeruginosa*, and *S. aureus*) by agar well diffusion [49].

The antimicrobial action of Se-NPs was reliant concentration As the Se-NPs succeeded to displayed a high antimicrobial action against *B. subtilis* and *E. coli* at 100 µg mL⁻¹ followed by concentrations of 75, 50, and 25 µg mL⁻¹ [50]. In the same vein the greatest antibacterial activity of Se-NPs against *B. subtilis*, *S. aureus*, *P. aeruginosa*, *E. coli*, *C. albicans*, *C. tropicalis*, *C. glabrata*, and *C. parapsilosis* was gotten at the concentration of 400 µg mL⁻¹ and lessened at low at 200 µg mL⁻¹ [51].

Blood parameters can reveal important information about an organism's internal health. The current study discovered that Se-NP supplementation enhanced the blood indices of *O. niloticus* fingerlings. A prior study demonstrated enhanced haematology in juvenile common carp, indicating a better health profile for fish fed with Se-enriched diets [52]. Similarly, it was found that Se-NPs supplementation raised all hematological markers, which coincide with our results [53]. Additionally, Nile tilapia fed dietary Se-NPs (0.8mg/kg) had higher levels of HCT, Hb, RBCs, and WBCs than tilapia fed bulk-selenium or a control diet [54]. These findings are related to the regulation of metabolic rate induced by Se-NPs, as the higher the number of RBCs and Hb content, the greater the availability of oxygen in human tissues [55]. Selenium's potent antioxidant activity [56] may increase RBC membrane balance and life span by protecting them from damaging oxygen-free radicals that cause anaemia, avoiding membrane rupture, cell hemolysis, and degeneration, and ensuring erythrocyte health and integrity [57].

A. hydrophila infection resulted in a considerable decrease in RBCs count, Hb concentration, and Hct% which can be attributable to the hemolytic activity of β-hemolysin. This hemolysin can cause RBC lysis, resulting in anemia, as previously documented [58].

The substantial fall in haemoglobin levels reported after the infection could be owing to a higher rate of RBC breakdown by pathogenic bacteria and/or a slower rate of RBC production [59]. Hemoglobin content falls due to RBC enlargement and inadequate haemoglobin mobilization by the spleen and other haematopoiesis organs [30]. In the

current investigation, the considerable fall in hemoglobin following fish infection could be due to severe anaemia. The anaemic reaction could be caused by the loss of intestinal cells that produce vitamin B12, which is utilized in the formation of the haemoglobin portion of red cells, haemodilution, or disturbance in erythrocyte development [60]. Furthermore, septicemia caused by the infection may result in RBC lysis due to the action of bacterial enterotoxins. The observed variations in mean corpuscular volume (MCV) indicate erythrocyte swelling, which reflects macrocytic anemia. The increase in MCV could be attributable to erythrocyte enlargement caused by hypoxic circumstances, disturbed water balance, or osmotic stress [61].

In the current study, WBCs revealed an increasing trend in the Se-NPs supplemented group (G 2), which could be attributed to the fish's enhanced health profile, which includes an increased immune response and various cell-mediated physiological activities. These alterations were attributed to a powerful antioxidant component of selenium that increased the lifespan of blood cells [62]. In fact, diets enriched with Se-NPs significantly enhanced fish cell-mediated immunity by long-term Se supplementation, as seen by rising WBC numbers in fish [63]. The current study found that the number of neutrophils, lymphocytes, and monocytes increased considerably ($P < 0.05$) in group 2 compared to the control. The current findings are consistent with those of another researcher, who demonstrated that in *O. niloticus*, selenium alters and controls the expression of immune-regulated selenoprotein [18].

Leukocytes play a protective function against pathogenic bacterial infections by activating the nonspecific defense system. An increase in leukocyte cells promotes faster recovery from *A. hydrophila* infection [64]. Total leukocytes increased following *A. hydrophila* infection, indicating a fish body defense response against existing antigens [65]. This finding is consistent with prior research, which found that WBC levels rose in tilapia infected with *A. hydrophila* [58]. WBCs are the primary component of the body's front lines of defense, and their numbers rapidly increase when an infection occurs. Increased WBC count in sick fish may serve as a protective barrier against pathogenic infestation [66].

Fish injected with *A. hydrophila* had considerably fewer lymphocytes than those not infected ($P < 0.05$). Decreases in lymphocytes following *A. hydrophila* infection of the Nile tilapia has been linked to cell re-trafficking to lymphoid tissues, resulting in their clearance from the blood stream [67].

Tilapia's monocyte and neutrophil levels increased during the challenge test. This is the fish body's attempt to absorb antigens that enter it, as

seen by an increase in the phagocytosis index following infection with *A. hydrophila*. Neutrophils and monocytes are important components of the fish's innate immune system, especially in the elimination of foreign substances [68].

Neutrophils and monocytes are professional phagocytes that consume and destroy infections by phagocytosis [69]. They are engaged in the identification and clearance of foreign entities such as pathogens and cellular debris, which helps to maintain tissue integrity and homeostasis [70]. Fish have a sophisticated innate immune response, with neutrophils and macrophages playing critical roles in pathogen identification, killing, and the initiation of adaptive immune responses [71].

The increase in Hb concentration and RBC count in group 4 treated with Se-NPs could be attributed to Selenium antioxidant activity, which protects the Hb molecule and the erythrocyte membrane from oxidative damage while also restoring erythropoiesis [56]. The increase in leukocytic and lymphocytic counts may be attributed to the immunostimulant effect of Se-NPs, which promotes leukocyte proliferation, the antibacterial efficacy of selenium in neutralizing infection, and the preservation of the leukocyte redox status [72].

The current study found that *Aeromonas* infection in Nile tilapia caused significant changes in liver function biomarkers, as revealed by significantly higher serum ALT and AST levels in the infected group compared to the control and Se-NPs treated groups. This rise reflects hepatic damage and leaking of these enzymes into the bloodstream due to structural and functional degradation of the liver tissue, a common reaction to bacterial septicemia and endotoxin exposure [73]. The observed non-significant reduction in ALT, along with a significant decline in AST activity in group 4, suggests a hepatoprotective role of nano-selenium, likely mediated through its antioxidant capacity and free radical scavenging properties [9].

Similarly, serum ALP and GGT activities were significantly raised in infected fish, indicating cholestasis and biliary epithelial damage linked with *Aeromonas*-induced hepatic injury [74]. Nano-selenium treatment considerably lowered both values, indicating enhanced hepatobiliary function and membrane stabilization, similar with prior studies on selenium-mediated protection in fish challenged with bacterial pathogens [75].

The protein profile analysis revealed that *Aeromonas* infection dramatically lowered total protein, albumin, and globulin levels, indicating poor protein synthesis and potential protein loss by tissue exudation during systemic inflammation [62]. Globulin reduction also suggests a suppressed humoral immune response. A decreased humoral immune response is also shown by a decrease in

globulin. Although values were still lower than those of uninfected controls, the addition of nano-selenium to infected fish greatly improved these parameters when compared to the infected group. The enhancement of immune cell function and protein production by selenium may be the cause of this improvement [76].

In terms of kidney function, fish infected with *Aeromonas* had higher levels of serum urea, creatinine, and uric acid, indicating renal impairment and decreased clearance capacity, most likely as a result of glomerular and tubular damage brought on by bacterial toxin [77]. Urea, creatinine, and uric acid levels were all markedly reduced after receiving nano-selenium, indicating nephroprotection via anti-inflammatory and antioxidant properties [78].

Antioxidant research revealed that infected fish had considerably lower levels of SOD, CAT, and GPx activities, indicating severe oxidative stress. This is consistent with the bacterial infection's increased production of reactive oxygen species (ROS), which overwhelms the body's built-in antioxidant defenses [4]. SOD, CAT, and GPx activity were all markedly restored by nano-selenium therapy. These results are consistent with the known function of selenium as a crucial part of selenium-dependent enzymes such as GPx, which aid in ROS detoxification and oxidative damage prevention [10]. Moreover, selenium nanoparticle-supplemented diets for European sea bass (*Dicentrarchus labrax*) reduce oxidative stress and increase the activity of oxidative enzymes [79]. Furthermore, when fish were exposed to pathogens that cause disease, the addition of selenium nanoparticles enhanced their cellular defenses against oxidative stress and increased their antioxidant defenses [80].

Selenium is a structural and functional component of GPx as well as a cofactor [81]. As a result, GPx activity may serve as a possible indicator of dietary Se supplementation and aid in the reduction of oxygen free radicals and detoxification of lipid hydroperoxides [82]. Furthermore, the use of dietary Se-NPs that promote glutathione formation could greatly reduce oxidative stress [83].

According to some studies, nanomaterials have gained popularity in fish farming because of their beneficial qualities (disease resistance and immunostimulant) and lower toxicity when compared to their organic and/or inorganic sources [84]. Furthermore, Se-NPs' immune-stimulant, antioxidant, bioavailable, and low toxicity properties may lessen the bacterial challenge in *O. niloticus* [85].

One of the key defense processes in the fish immune system is phagocytosis, which occurs when macrophages and neutrophils ingest and remove pathogens. Phagocytes are white blood cells that help the innate immune system engulf and remove

invading particles. Phagocytes' ability to affect other immune components in the blood and tissue allows them to act as a link between innate and adaptive immunity. Fish phagocytes are mostly composed of macrophages, monocytes, granulocytes, and dendritic cells, but antibody-producing cells (B cells) have also been shown to have bactericidal and phagocytosis activities [86]. Fish phagocytes contribute to the engulfment of harmful bacteria, which reduces bacterial pathogenic movement, destroys pathogenic bacterial cells, and thereby reduces bacterial infection in the host [887].

Selenium is a component of selenoprotein, which can boost serum protein production, therefore taking Se-NPs as a dietary supplement may improve non-specific immunity [88].

The phagocytic index and phagocytic percentage were observed to significantly rise with Se-NPs supplementation in the present investigation. Our findings are supported by previous work [89], which reported an increased lysozyme level, pro-inflammatory cytokines (TNF- α and IL-1 β), and macrophage activity after supplementing Nile tilapia with nano-Se. Dietary Se nanoparticles (1-2 mg/kg) improved serum lysozyme and respiratory burst activity in Nile tilapia. Additionally, Nile tilapia given dietary Se nanoparticles (1-2 mg/mg) demonstrated increased phagocytic and lysozyme activity, phagocytic index, globulin, and immunoglobulin M, according to [90]. Furthermore, the total serum protein globulin, phagocytic index, phagocytic, and lysozyme activity of European seabass were increased by dietary Se nanoparticles (0.5-1 mg/kg) [91].

Bacterial infections such as *A. hydrophila* can trigger an increase in phagocytic activity as a first line of defense against pathogen transmission in the fish body. Detection of pathogen-associated molecular patterns (PAMPs) by immune cell receptors results in the production of proinflammatory cytokines and the activation of phagocytes, which is the cause of this increase [92].

Conclusion

Overall, the findings indicate that nano-selenium supplementation in *Aeromonas*-infected Nile tilapia reduces hepatic and renal damage, improves protein metabolism, and boosts antioxidant levels. This protective benefit is most likely due to selenium's dual role in improving immunological function and combating oxidative stress, which supports organ function and overall health in diseased fish on the bacteriological level (Se NPs) exhibited excellent antibacterial activity for MDR *A. hydrophila* isolates, so it can be considered as an appropriate and environment-friendly therapeutic measure. Finally, it is recommended that additional research be conducted on the antimicrobial susceptibility tests for *Aeromonas*.

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Declaration of Conflict of Interest

The authors declare that there is no conflict of interest.

Ethical of approval

This work reviewed and approved by the ZUIACUC committee at the Faculty of Veterinary Medicine, Zagazig University, Egypt. Approval number: ZU IACUC/2/F/20/2025.

TABLE 1. Oligonucleotide primer sequences, its product size and cycling conditions during PCR [22].

Gene	Primers sequences	Product size (bp)	Amplification (35 cycles)					Reference
			Primary Denaturation	Secondary denaturation	Annealing	Extension	Final extension	
<i>A.hydrophila</i> 16S rRNA	GAAAGGTTGATGCCTAATACGTA CGTGCTGGCAACAAAGGACAG	685	94°C 5 min.	94°C 30 sec.	50°C 40 sec.	72°C 45 sec.	72°C 10 min.	22

TABLE 2. Breakpoints of the disk diffusion methods used to determine the antimicrobial susceptibility of *Aeromonas* spp.

Antimicrobial agent	Symbol	Disc potency (µg)	Resistant S	Intermediate I	Sensitive R
Doxycyclin	DO	30	≥14	11-13	≤10
Trimethoprim-sulfa- methoxazole	SXT	1.25/23.75	≥19	-	<16
Ciprofloxacin	CIP	5	≥27	-	<24
Chloramphenicol	C	30	≥18	13-17	12
Gentamicin	CN	10	≥18	16-17	<16
Neomycin	N	30	≥23	13-16	<13
Streptomycin	S	10	≥15	12-14	≤11
Erythromycin	E	15	-	-	-
Kanamycin	K	30	≥18	14-17	≤13
Amoxicillin	AX	25	≥19	16-18	<16
Nalidixic acid	NA	30	≥20	15-19	<15
Colistin	CT	10	Not available	Not available	Not available
Cefepime	FEP	30	≥27	-	<24

zone diameter breakpoints, according to the Committee for Antibiogram of the French Society of Microbiology [26]

TABLE 3. Antimicrobial sensitivity test results of *A. hydrophila* obtained with disc diffusion method

Antimicrobial agent	Symbol	Disc potency (µg/ Disc)	Isolates %		
			S	I	R
Doxycyclin	DO	30	-	-	100
Trimethoprim-sulfa- methoxazole	SXT	1.25 /23.75	-	-	100
Ciprofloxacin	CIP	5	-	-	100
Chloramphenicol	C	30	-	-	100
Neomycin	N	30	--	25	75
Streptomycin	S	10	-	-	100
Erythromycin	E	15	-	-	100
Kanamycin	K	30	-	-	100
Amoxicillin	AX	25	-	-	100
Nalidixic acid	NA	30	-	-	100
Colistin	CT	10	-	-	100
Cefepime	FEP	30	-	-	100
Gentamicin	CN	10	33.3	-	66.6

TABLE 4. Effect of Se-NPs supplementation on hematological variables of *O. niloticus* challenged with *A. hydrophila*

Group Parameters	Group1(Control)	Group 2 (Se- NPs)	Group 3 (Infected control)	Group 4
RBCs X 10 ⁶ /cmm	2.78 ^b ± 0.10	3.36 ^a ± 0.15	1.72 ^d ± 0.14	2.14 ^c ± 0.09
Hb g/dl	6.28 ^b ± 0.21	7.43 ^a ± 0.24	5.02 ^c ± 0.16	5.84 ^b ± 0.19
Hct %	26.46 ^b ± 0.55	29.80 ^a ± 0.44	22.64 ^c ± 0.38	24.56 ^b ± 0.62
MCV fl	95.18 ^c ± 2.88	88.70 ^c ± 2.63	131.62 ^a ± 4.62	114.77 ^b ± 3.11
MCH Pg	22.59 ^b ± 0.57	22.11 ^b ± 0.45	29.18 ^a ± 0.59	27.28 ^a ± 0.61
MCHC g/dl	23.73 ^a ± 0.60	24.93 ^a ± 0.63	22.17 ^a ± 0.57	23.77 ^a ± 0.65

* Values represent means ± standard error (n = 5). Means with distinct small letters in the same row show significant differences (P < 0.05).

TABLE 5. Effect of Se-NPs supplementation on leukogram of *O. niloticus* challenged with *A. hydrophila*

Group Parameters	Group 1 Control	Group 2 Se-NPs	Group 3 Infected control	Group 4
WBCs X10 ³ /cmm	12.83 ^c ± 0.41	14.45 ^b ± 0.36	16.88 ^a ± 0.42	14.92 ^b ± 0.40
Neutrophils X10 ³ /cmm	3.43 ^d ± 0.11	3.95 ^c ± 0.19	7.96 ^a ± 0.32	5.49 ^b ± 0.24
Lymphocytes %X10 ³ /cmm	6.82 ^b ± 0.22	7.66 ^a ± 0.26	5.47 ^d ± 0.21	6.24 ^c ± 0.16
Monocytes X10 ³ /cmm	1.63 ^c ± 0.04	1.85 ^b ± 0.05	2.36 ^a ± 0.07	2.15 ^a ± 0.08
Eosinophils X10 ³ /cmm	0.69 ^a ± 0.03	0.71 ^a ± 0.04	0.79 ^a ± 0.07	0.73 ^a ± 0.05
Basophils X10 ³ /cmm	0.26 ^a ± 0.03	0.28 ^a ± 0.02	0.30 ^a ± 0.03	0.31 ^a ± 0.02

* Values represent means ± standard error (n = 5). Means with distinct small letters in the same row show significant differences (P < 0.05).

TABLE 6. Effect of Se-NPs supplementation on biochemical parameters of *O. niloticus* challenged with *A. hydrophila*

Group Parameters	Group 1 Control	Group 2 Se-NPs	Group 3 Infected control	Group 4 Aeromonas+ nano
ALT U/L	9.21 ^b ± 0.16	9.51 ^b ± 0.37	18.20 ^a ± 2.534	14.77 ^a ± 0.99
AST U/L	101.91 ^c ± 1.22	102.96 ^c ± 0.90	152.10 ^a ± 1.154	116.23 ^b ± 1.58
ALK U/L	11.10 ^c ± 0.53	12.20 ^c ± 0.57	19.20 ^a ± 0.55	16.26 ^b ± 0.56
GGT U/L	1.17 ^c ± 0.117	1.16 ^c ± .031	2.93 ^a ± 0.199	1.59 ^b ± 0.005
total protein g/dl	3.42 ^a ± 0.15	3.48 ^a ± 0.06	2.23 ^c ± 0.060	2.92 ^b ± 0.035
Albumin g/dl	1.92 ^a ± 0.14	1.95 ^a ± .036	1.28 ^c ± 0.044	1.60 ^b ± 0.057
Globulin g/dl	1.50 ^a ± 0.048	1.53 ^a ± 0.046	0.95 ^c ± 0.018	1.32 ^b ± 0.014
Urea Mg/dl	0.88 ^c ± 0.020	0.84 ^c ± 0.026	1.72 ^a ± 0.133	1.20 ^b ± 0.057
Creatinine Mg/dl	0.28 ^c ± 0.014	0.26 ^c ± 0.015	0.57 ^a ± 0.018	0.40 ^b ± 0.005
Uric Acid Mg/dl	0.71 ^c ± .026	0.70 ^c ± 0.024	0.99 ^a ± 0.017	0.82 ^b ± 0.029

* Values represent means ± standard error (n = 5). Means with distinct small letters in the same row show significant differences (P < 0.05).

TABLE 7. Effect of Se-NPs supplementation on oxidative stress responses, phagocytic % and phagocytic index of *O. niloticus* challenged with *A. hydrophila*

Group Parameters	Group 1 Control	Group Se-NPs	Group 3 Infected control	Group 4
SOD (U/mg protein)	51.29 ^b ± 0.55	88.6 ^a ± 0.32	38.6 ^d ± 0.32	47.63 ^c ± 1.06
Cat (U/mg protein)	2.62 ^b ± .14518	5.49 ^a ± 0.44	1.04 ^d ± 0.41	2.09 ^c ± 2.81
Gpx (U/mg protein)	61.10 ^b ± 0.52	99.61 ^a ± 1.57	45.76 ^d ± 1.2	55.26 ^c ± 2.59
Phagocytic %	49.68 ^c ± 0.82	54.82 ^b ± 0.69	61.14 ^a ± 0.73	60.25 ^a ± 0.98
Phagocytic index	1.83 ^c ± 0.13	2.51 ^b ± 0.16	3.94 ^a ± 0.14	3.76 ^a ± 0.18

* Values represent means ± standard error (n = 5). Means with distinct small letters in the same row show significant differences (P < 0.05).



Fig.1. *A. hydrophila* on Aeromonas Base medium: Opaque green colonies with a darker center

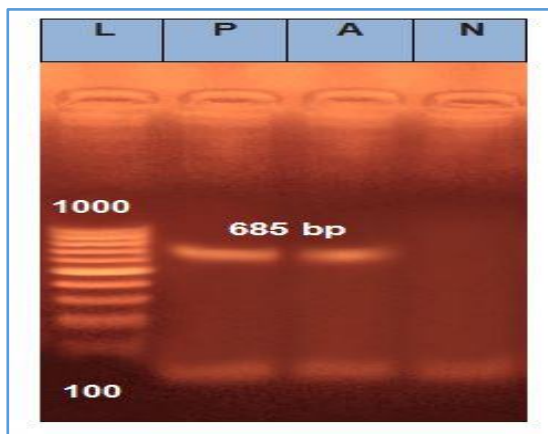


Fig.2. uniplex: 16s rRNA gene specific for *A. hydrophila* (685 bp)

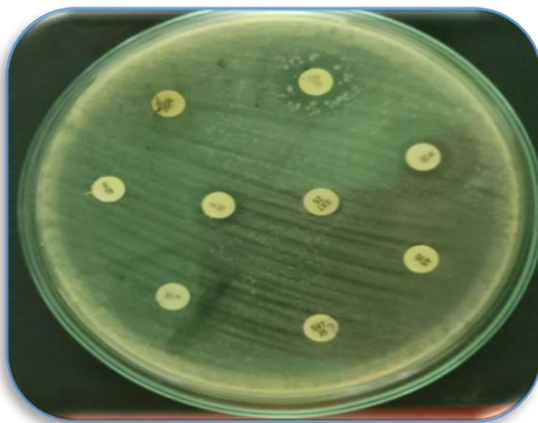


Fig. 3_a. *A. hydrophila* showing multidrug resistance via disc diffusion sensitivity test

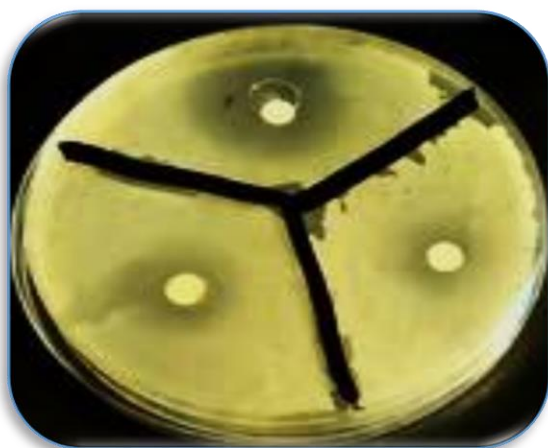


Fig. 3. disc diffusion technique and zone of inhibition after exposure of *A. hydrophila* to Se NPs showing inhibition zones ranging from 11mm: 28mm.

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دراسة تأثير جزيئات السيلينيوم النانوية على مقاومة الايرومونات في أسماك البلطي النيلي

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الملخص

أحد أهم مسببات المرضية التي تصيب أسماك المياه الدافئة، *Aeromonas hydrophila* تعد بكتيريا وخاصة أسماك البلطي النيلي، حيث تؤدي إلى ظهور العديد من الأمراض البكتيرية. هدفت هذه الدراسة إلى *hydrophila* بحث القيود المرتبطة باستخدام المضادات الحيوية في علاج العدوى الناجمة عن (كخيار علاجي بديل. أجريت الدراسة في Se-NPs) بالإضافة إلى تقييم فعالية جسيمات السيلينيوم النانوية محافظة الشرقية بجمهورية مصر العربية، حيث أظهرت النتائج انتشاراً مرتفعاً لبكتيريا بنسبة 92% من العزلات البكتيرية، وجميعها أبدت مقاومة متعددة للمضادات الحيوية شملت 11 من أصل 13 مضاداً حيوياً تم اختبارها. تم تنفيذ تجربة تغذية استمرت 35 يوماً على 120 من أسماك البلطي النيلي، بجزيئات قُسمت إلى أربع مجموعات: مجموعة ضابطة سالبة، مجموعة تغذت على علائق أساسية مدعمة ومجموعة مصابه تغذت علي جزيئات *A. hydrophila* النانو سليينيوم ومجموعة ضابطة موجبة مصابه ب النانو سليينيوم. وظهرت النتائج ان العدوي أحدثت تغيرات في المؤشرات الدموية والبيوكيميائية، تمثلت في ارتفاع مستويات كل من اليوريا والكرياتينين وحمض (ALT, AST, ALP, GGT) إنزيمات الكبد البولييك، إلى جانب انخفاض البروتين الكلي، الألبومين، الجلوبيولين، وإنزيمات مضادات الأكسدة. وعلى النقيض، فإن إضافة جزيئات النانو سليينيوم إلى العليقة حسنت بصورة ملحوظة هذه المعايير، من خلال تعزيز النشاط المضاد للأكسدة وتنشيط الاستجابة المناعية في الأسماك. علاوة على ذلك، أسهمت جزيئات النانو سليينيوم في التخفيف من التأثيرات المرضية الناتجة عن العدوى، مما يشير إلى إمكان استخدامها كخيار علاجي واعد للسيطرة على الأمراض البكتيرية في أنظمة الاستزراع السمكي.

الكلمات الدالة: سمك البلطي النيلي، جسيمات السيلينيوم النانوية، بكتيريا إيرومونات هيدروفيليا، أمراض الدم. مضادات الأكسدة، نسبة البلمعة.