



Determination of antibiotic resistance pattern in *Aeromonas hydrophila* isolated from Reared *Oncorhynchus mykiss* in Tabriz city, Iran

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ABSTRACT

Objective: *Aeromonas hydrophila* is a Gram negative, positive oxidase, anaerobic, and opportunistic bacteria that, under certain conditions, become a pathogen (in humans and fish). This bacterium causes toxin and host infection in which different antibiotic resistance in isolated strains has been reported in different regions of the world. The aim of this study was to determine the prevalence of this bacterium and its susceptibility to common antibiotics in Tabriz city.

Materials and Methods: 50 samples from 5 Reared *Oncorhynchus mykiss* farms in Tabriz city (For each farm, 10 numbers) were randomly assigned to suspected fish to the disease.

Results: By using biochemical tests, 14 samples (28%) from 2 Fish farms (40%) were identified as *A. hydrophila*. Antibiogram for these specimens showed that the bacterium had the highest resistance to vancomycin (100 %) and clindamycin (92.8%) antibiotics, and has the most sensitivity to the antibiotics of sultrim, tetracycline and oxytetracycline with 71.4%.

Discussion: Considering the different antibiotic resistance pattern in this study and other similar studies, the necessity of examining the pattern of resistance in each region seems necessary.

Keywords: *Aeromonas hydrophila*, *Oncorhynchus mykiss*, Antibiotic resistance

INTRODUCTION

The *Aeromonas* spp. were originally located in the *Vibrionaceae* family, but after genetic studies, it was found that they are not closely related to the vibrio. Thus, the *Aeromonas* spp. are located in a new family called *Aeromonadaceae* (Janda and Abbott, 2010; Ceylan et al., 2009; Gülaydın et al., 2018). Pathogenicity of *Aeromonas* species is very common in drinking water samples and types of food samples, especially marine foods. Today, marine products for humans are one of the most important food products. These products can easily be contaminated with a variety of pathogens (Puthucheary et al., 2011; Villari et al., 2003; Yogananth et al., 2009). *Aeromonas hydrophila* is an

opportunistic pathogen, which can host a wide range of hosts, including various species of fish such as *Cyprinus carpio*, catfish, Tilapia, Eels, Goldfish, Trout and humans (Xia et al., 2004; Zhou et al., 2010). Today, the presence of the bacterium *A. hydrophila* in the aquatic ecosystem is well documented. This bacterium is part of the aquatic intestinal flora, which can be pathogenic when caused by tension (Yogananth et al., 2009). *A. hydrophila* causes many complications and diseases in fish, including Hemorrhagic Septicemia, Dark skin, skin ulcers, Pop-Eye and asymptomatic septicemia. This bacterium often causes irreparable damages in farmed fish populations and infection control depends on eliminating the predisposing

factors of the infection (Khushiramani et al., 2007; Jayavignesh et al., 2011). Recent studies indicate that, based on the severity of infection and the age of breeding fish with *A. hydrophila*, different clinical symptoms and even mortality are expected (Esmaeli and Peighan, 1997; Vega-Sánchez et al., 2014). In the last few decades, the widespread use of antibiotics in the treatment of human, animal, agricultural and Fishery infections has increased the presence of these compounds in the environment and, consequently, the resistance of some bacteria (Yours et al., 2007). Today, trout (*Oncorhynchus mykiss*) is known as one of the most important fish species in Most of the cold-water fish farms, in most parts of the world (Tafi et al., 2014). Considering that trout (*Oncorhynchus mykiss*) is a diet of human societies, the aim of this study was to determine the antibiotic resistance pattern in *A. hydrophila* isolated from trout in Tabriz city.

MATERIALS and METHODS

Sampling and biochemical identification of isolates:

In this experimental study, 50 trout samples with suspected symptoms were collected from 5 farms in Tabriz city (10 fish per farm) at a 20-day interval from June to September 2018 and in suitable conditions and in the transport box of fish, with appropriate oxygenation was transferred to the microbiological laboratory. The Symptoms in all suspected fish included: abdominal distension, Inflammation of the Anus, Pop-Eye and Dark skin. All field conditions such as water factors (temperature and oxygen), food (type and physical and chemical characteristics) and fish (apparent observations) were recorded before the experiment was performed. Prior to sampling, each piece of fish was killed by cutting the spinal cord and then weighed. The body surface of the fish was washed with 70 ° C ethanol and the liver and Kidney and all of them were isolated from the abdominal area under sterile conditions and were washed twice with sterile physiology serum. By inserting the Inoculation needle into the tissues, the specimens were transferred to a blood-agar culture medium and incubated for 24 hours at 22 ° C. Suspicious colonies (white to gray, convex and semi-transparent colonies) based on biochemical tests including mobility, Gram stain, oxidase and catalase tests, indole, methyl red, citrate, nitrate recovery, sulfate production, KOH Identification and examples with features Gram-negative, Catalase and Oxidase-positive properties were

transferred to Shotts-Rimler medium (Shotts and Rimler, 1973) as a selective medium of *A. hydrophila*.

Antibiotic susceptibility testing:

To determine the antimicrobial resistance pattern, disk diffusion method was used in Muller Hinton Agar (Merck, Germany) culture media (Bauer et al., 1966). Antibiotic disks include Vancomycin (30 µg), Ciprofloxacin (25 µg), Amoxicillin (30 µg), Sultrim (10 µg), Tetracycline (30 µg), Streptomycin (20 µg), Gentamicin (15 µg), Cefotaxime (25 µg), Nalidixic Acid (30 µg), Oxytetracycline (20 µg), Chloramphenicol (20 µg) and Clindamycin (30 µg) from the company PADTAN TEB(Iran) were used. Interpretation of antibiogram results was performed according to the CSLI (CLSI, 2009) standard.

RESULTS

Out of 50 samples taken from 5 trout farms with suspected symptoms, 14 specimens (28%) from 2 farms (40%) were identified as positive *A. hydrophila* specimens. By analyzing the results of the antibiogram, 14 isolates of *A. hydrophila* and the standard strain of *A. hydrophila* showed that the highest resistance among the isolates was related to vancomycin antibiotics (100%) and clindamycin (92.8%) and the least resistance in The isolates were associated with gentamicin antibiotics (7.2%) and ciprofloxacin (14.2%), and the most susceptible to sultrim, tetracycline and oxytetracycline antibiotics was 71.4%.

Table 1. Results of study on determination of microbial susceptibility of isolated strains of *A. hydrophila* (%)

Standard antibiotics	R	I	S
Vancomycin	100	0	0
Ciprofloxacin	14.3	35.7	50
Amoxicillin	28.6	50	21.4
Sultrim*	21.4	7.2	71.4
Tetracycline	21.4	7.2	71.4
Streptomycin	71.4	0	28.6
Gentamicin	7.2	42.8	50
Cefotaxime	0	42.8	57.2
Nalidixic Acid	0	50	50
Oxytetracycline	21.4	7.2	71.4
Chloramphenicol	0	57.2	42.8
Clindamycin	92.8	0	7.2

*: Trimethoprim – Sulphadiazine; R: Resistant; I: Intermediate; S: Sensitive;

DISCUSSION and CONCLUSION

One of the causes of heavy losses in fish farms is bacterial infection, which causes severe economic damage to the aquaculture industry annually (Nielsen et al., 2001). Among these, the *A. hydrophila* bacteria is one of the most important opportunistic bacteria in aquatic organisms, which causes Hemorrhagic Septicemia and high mortality in aquaculture system (Casiano et al., 2010). Although other factors such as viral infection, nutrition, environmental factors also contribute to this mortality. But some studies have shown the presence of *Aeromonas* in these high-profile casualties (Ahangarzadeh et al., 2017). In this study, 50 specimens from 5 trout farms with suspicious symptoms were identified as 14 specimens (28%) from 2 farms (40%) as the positive specimen of *A. hydrophila*. Fadaeifard, of the 60 samples taken from 10 farms, identified 9 samples (15%) of *A. hydrophila* Positive (Fadaeifard, 2014). Igbal et al. (1999) examined various species of *Aeromonas*, which showed that the high percentage of *Aeromonas* isolated to tetracycline is susceptible. Which was consistent with the findings of the present study. Arzani et al. (2016), by investigation the antibiotic susceptibility of *A. hydrophila* bacteria in Tonekabon fish farming pools showed that *A. hydrophila* isolates were 100% sensitive to antibiotics tetracycline, gentamicin, kanamycin, norfloxacin, ofloxacin, amikacin, nalidixic acid, azithromycin and ciprofloxacin, and 100% resistant to oxacillin, meticillin, erythromycin, ampicillin and amoxicillin. Bakhtiari et al. (2017) in Khuzestan province showed the highest resistance of *A. hydrophila* to vancomycin and clindamycin (90%) streptomycin (70%) and the highest susceptibility to Sultrim, tetracycline and oxytetracycline (75%). Which are consistent with the findings of this study. Hosseinzadeh and Tukmechi (2016) with the study of *A. hydrophila* isolated from common carp, showed that the bacteria were most susceptible to ciprofloxacin (84.11%), oxytetracycline (59.44%), gentamicin (59.22%) and streptomycin (41.06%). Sarder et al. (2016) by Examined antibiotic susceptibility of *A. hydrophila* isolated from freshwater fish Showed that the bacteria had the highest susceptibility to gentamicin antibiotics (100%), ciprofloxacin (93.53%), chloramphenicol (81.25%) and the highest resistance to amoxicillin (93.75%). Bektas and Iscimen (2005) in Turkey, by studying *A. hydrophila*, showed that the bacterium is susceptible to chloramphenicol, tetracycline and oxytetracycline antibiotics and resistant to amoxicillin

antibiotics. The antibacterial function of tetracycline is by through the disruption of the translation of the Aerolysin gene and thus inhibit protein synthesis in the bacterial ribosome, thus, by stopping the production of proteins in *A. hydrophila*, these bacteria are condemned to death (Pakravan and Akbarzadeh, 2017). Revina et al. (2017) examined antibiotic resistance patterns in *A. hydrophila* isolates, which showed that all isolates were resistant to amoxicillin, ampicillin, cefalexin and erythromycin and had the lowest resistance to gentamicin (4.4%), Streptomycin (10%). Saavedra et al., (2004) determined that *A. hydrophila* isolated from rainbow trout showed the highest resistance to antibiotic amoxicillin (88%) and the most sensitive to cefotaxime (88%). The findings of this study have differences with similar studies. In order to justify this difference, as many studies have shown, a bacteria of one species may have several subtypes. Also, the acuity of these bacteria in different sources of water may vary. Therefore, the lack of specific information to describe the actuality of organisms that have been used in laboratory evaluation of the antimicrobial properties of bacteria in fish breeding or water resources can be considered as one of the problems. This study provides basic information on resistance to common antibiotics in *A. hydrophila* isolates from trout (*Oncorhynchus mykiss*) farms in Tabriz city. The findings of this study indicate that *A. hydrophila* bacteria is responsible for causing septicemia in trout (*Oncorhynchus mykiss*). It can also be said that if the bacteria enter the human food chain through infected fish, there is a risk of human bloody diarrhea. On the other hand, it can be concluded that the probability of a horizontal gene transfer is one of the determinants of antibiotic resistance in these isolates and excessive consumption of antibiotics in aquaculture has led to the emergence of resistant strains in *A. hydrophila*.

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