



Antimicrobial susceptibility profile of *Edwardsiella tarda* from farmed fish in Ilorin Metropolis, Kwara State Nigeria

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Abstract

This study determined the susceptibility profile and multiple antibiotic resistance (MAR) index of *Edwardsiella tarda* obtained from farmed fish in Ilorin Metropolis, Kwara State, Nigeria. 30 isolates were obtained from 30 catfish samples from the major fish market in Ilorin Metropolis. It was observed that the isolates showed resistance to tetracycline (100%), ampicillin (100%), cefotaxime (100%) and ciprofloxacin (97%). *Edwardsiella tarda* isolates were entirely susceptible to Gentamycin (100%) while also showing varying levels of susceptibility to Enfloxacin (77%), Norfloxacin (67%), Sulphamethoxazole (50%), Composulphonamides (23%) and Chloramphenicol (10%). The isolates showed an intermediate range of susceptibility to Chloramphenicol (90%), Composulphonamides (77%), Sulphamethoxazole (50%) and Ciprofloxacin (3%). Analysis of the MAR index of isolates revealed that 97% of the isolates had a MAR index of 0.4, with only 3% of the isolates showing a multiple antibiotic resistance index of 0.3, which is an indication that farmed fishes in the Ilorin metropolis had a high risk of exposure to antibiotics.

Keywords: Antimicrobial, Antibiotics, Bacteria, Resistance, Intermediate, Susceptibility, Susceptible, Multiple antibiotic resistance, Isolates, *Edwardsiella tarda*

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Introduction

Antibiotic abuse and uncontrolled use in the fish farming industry have significantly increased the amount of resistant microorganisms in the world including Nigeria. These antimicrobial practices have been widely used in fish farming industry exclusively for therapeutic or preventative purposes (Manyi-Loh *et al.*, 2018). However, these antibiotics are typically used to boost and speed up fish growth and for effective feed efficiency in the fish (Muhammad *et al.*, 2020). Antibiotics are widely used in Nigerian aquaculture and livestock husbandry. All antibiotic resistance in microbes has a genetic basis (Abu and Wondikom, 2018) since it is a defence mechanism that enables bacteria to survive in an unfavourable environment (Gang *et al.*, 2015). Antibiotic resistance represents one of the most significant current threats to public health and is predicted to overtake cancer as a cause of death by 2050. Nevertheless, little or no attention has been paid to the potential use of antibiotics in aquaculture industries. In addition to the transfer of antibiotic-resistant microorganisms and their genes through consuming contaminated fish, there is a significant risk of environmental contamination due to using medicated feeds.

Antibiotic resistance has been reported in catfish, shrimp, eels, and aquaculture environments. According to some accounts, resistance develops shortly after using antibacterial medications to treat infections, reducing their effectiveness in managing bacterial

fish diseases. Antimicrobial usage must, therefore, be carefully regulated and monitored in aquaculture and our environment (Ma *et al.*, 2021) to evaluate their impact and subsequent risk to the ecosystem. *Edwardsiella tarda* is the causative agent of Edwardsiellosis in fish, characterised by systemic hemorrhagic septicemia, internal abscesses and skin lesions (Ewing *et al.*, 1965). The bacterium is opportunistic, and disease outbreaks occur when poor environmental conditions prevail, like overcrowding, poor water quality conditions, high organic content and high temperature (Mohanty and Sahoo, 2007).

This study investigated the occurrence of antimicrobial resistance of *Edwardsiella tarda* from farmed fish within Ilorin metropolis in Kwara State to identify high-risk sources of contamination and address the issue of emerging antibiotic resistance of *Edwardsiella tarda* bacteria from fish farming.

Materials and methods

Description of study area

The study area is within the Ilorin metropolis, Kwara State, and is characterised by various economic activities such as fish farming, trading, and fishing. It is also an important industrial and commercial area with many local manufacturing and trade companies.

Collection of fish samples

30 fish samples were collected from the major fish market in Ilorin Metropolis,

Kwara State. Fish samples were aseptically dissected using a dissecting kit.

*Isolation and identification of *Edwardsiella tarda**

- Swabs were taken aseptically from the intestine of the dissected fish samples and were streaked on Petri dish-containing Tryptone soy agar (LAB M Ltd., United Kingdom) that was prepared according to manufacturers' instructions.
- The plates were incubated at 37°C for 24 hours and were observed for bacterial growth and distinct colonies were further sub-cultured on freshly prepared tryptone soy agar to obtain a pure culture of the isolates.
- *Edwardsiella tarda* isolates were then identified using gram staining technique, morphology, motility, and biochemical reaction tests such as oxidase test, catalase test, lactose fermentation and glucose fermentation according to (Berger's Manual of Determinative Bacteriology, 7th Edition).
- Measurement of the zone of inhibition (zi), The zone of inhibition was measured using meter rule in millimeters (mm).

Antibiotic susceptibility testing

Antibiotic Susceptibility testing was carried out with a commercially available antimicrobial susceptibility test single disc from (Oxoid Ltd, Basingstoke, RG24 8PW, United Kingdom) Pure culture of *Edwardsiella tarda* was picked from Tryptone soy

broth using a sterile wire loop and transferred to tubes each containing 5 mL of sterile physiological saline. The suspension was vortexed and adjusted to match 0.5 McFarland turbidity standards. Sterile swab sticks were then dipped, rotated and pressed firmly on the tube walls above the culture to remove excess inoculums from the swabs. This was then evenly swabbed on the dried surface of Mueller-Hinton agar (Oxoid Ltd., England) plates ensuring even distribution of the bacterium. 10 different antimicrobial susceptibility single discs were placed on the plates using sterile forceps and allowed to undergo incubation at a temperature of 37°C for 18 to 24 hours, after which the diameter of the zone of inhibition was measured with a white transparent meter rule in millimetre. The antimicrobial susceptibility test was carried out using the agar disk diffusion method as described by Bauer *et al.* (1966). The methods described above and the interpretations of results were done using the Clinical Laboratory Standard Institute (CLSI, 2020) manual. The ten antibiotics that were used have the following concentrations: Enfloxacin (ENR) (5µg), Ampicillin (AMP) (10µg), Norfloxacin (NOR) (10µg), Gentamycin (CN) (10µg), Cefotaxime (CTX) (5µg), Tetracycline (TE) (10µg), Composulphonamides (S3) (30µg), Sulphamethoxazole (SXT) (5µg), Chloramphenicol (C) (30µg), Ciprofloxacin (CIP) (5µg).

Determination of antibiotics resistance index

In vitro sensitivity test was done on random selected isolates of *Edwardsiella tarda* using disc diffusion method according the method carried out by Bauer *et al.* (1966) using different antimicrobial agents: Enfloxacin (ENR) (5 μ g), Ampicillin (AMP) (10 μ g), Norfloxacin (NOR) (10 μ g), Gentamycin (CN) (10 μ g), Cefotaxime (CTX) (30 μ g), Tetracycline (TE) (10 μ g), Composulphonamides (S3) (30 μ g), Sulphamethoxazole (SXT) (25 μ g), Chloramphenicol (C) (30 μ g), Ciprofloxacin (CIP) (5 μ g). The diameter of the inhibition zone was measured and interpreted according to clinical and laboratory standards institute (CLSI, 2020) manual. Isolates that showed resistance to more than two different antibiotic groups were multiple drug-resistant (MDR) isolates.

Determination of multiple antibiotics resistance (MAR) index

Multiple antibiotics resistance (MAR) index is an analytical measure or technique of analysis introduced in 1983 by Krumpelman which helps to categorize or group different sources of high and low-risk contamination of bacterial isolates based on their frequency of antibiotic resistance (Osundiya *et al.*, 2013). Multiple Antibiotic Resistance (MAR) index for each strain was identified according to Singh *et al.* (2010), MAR index= No. of resistance (Isolates classified as intermediate were considered sensitive for MAR index) / Total No. of tested antibiotics. The multiple antibiotics index is a measure used to identify the

risk of contamination that poses a potential hazard to humans. However, multiple antibiotics resistance value greater than 0.2 is a signal that the bacterial isolates were from high-risk sources while bacterial isolates with multiple antibiotics resistance index less than 0.2 is an indicator of low-risk contamination (Nyandjou *et al.*, 2019).

Interpretation of the data obtained with clinical laboratory standard institute, CLSI (2020) Manual

The Clinical Laboratory Standard Institute Manual 2020 was used for the interpretation of the diameter (mm) of the zone of inhibition of the antimicrobials carried out during the experimental work (Table 1).

Statistical analysis

The data (diameter of inhibition zone) was measured and interpreted according to clinical and laboratory standards institute (CLSI, 2020) manual. The data obtained was analyzed using descriptive statistics with the aid of SPSS version 25.

Results

Susceptibility profile of Edwardsiella tarda Isolates (n = 30) from farmed fishes to 10 antibiotics

The data (diameter of inhibition zone) was measured and interpreted according to clinical and laboratory standards institute (CLSI, 2020) manual. The table below shows the susceptibility percentage of *Edwardsiella tarda* isolates tested with ten antibiotics, it can be inferred that all 30 isolates showed a

100% resistance to tetracycline, ampicillin and cefotaxime antibiotics while showing a 97% resistance to ciprofloxacin. Whereas the isolates showed a 0% resistance to Enfloxacin, Gentamycin, Norfloxacin, Chloramphenicol, Composulphonamides, and Sulphamethoxazole. Gentamycin was 100% susceptible to all the isolates while Enfloxacin, Norfloxacin, Sulphamethoxazole and Composulphonamides showed 77%, 67%, 50% and 23% susceptibility respectively. Chloramphenicol Composulphonamides, Sulphamethoxazole, Enfloxacin, Norfloxacin, and Ciprofloxacin also showed an intermediate range of 90%, 77%, 50%, 33%, 23% and 3% respectively (Table 2).

Table 1: Clinical Laboratory standard institute, CLSI (2020) manual.

Zone of Inhibition (mm)	Susceptible (S)	Intermediate (I)	Resistant (R)
Enfloxacin (ENR)	≥18	15 – 17	≤14
Gentamycin (CN)	≥15	13 – 14	≤12
Chloramphenicol (C)	≥18	13 – 17	≤12
Ciprofloxacin (CIP)	≥26	22 – 25	≤21
Tetracycline (TE)	≥15	12 – 14	≤11
Ampicillin (AMP)	≥17	14 – 16	≤13
Norfloxacin (NOR)	≥17	13 – 16	≤12
Cefotaxime (CTX)	≥26	23 – 25	≤22
Composulphonamides(S3)	≥17	13 – 16	≤12
Sulphamethoxazole (SXT)	≥16	11 – 15	≤10

Table 2: Susceptibility profile of *Edwardsiella tarda* Isolates (n = 30) from farmed fishes to 10 antibiotics.

Antimicrobial Agents	Disk Content	Susceptible Isolates (%)	Intermediate Isolates (%)	Resistant Isolates (%)
Enfloxacin (ENR)	5µg	23(77%)	7(23%)	0(0%)
Gentamycin (CN)	10µg	30(100%)	0(0%)	0(0%)
Chloramphenicol (C)	30µg	3(10%)	27(90%)	0(0%)
Ciprofloxacin (CIP)	5µg	0(0%)	1(3%)	29(97%)
Tetracycline (TE)	10µg	0(0%)	0(0%)	30(100%)
Ampicillin (AMP)	10µg	0(0%)	0(0%)	30(100%)
Norfloxacin (NOR)	10µg	20(67%)	10(33%)	0(0%)
Cefotaxime (CTX)	30µg	0(0%)	0(0%)	30(100%)
Composulphonamides (S3)	30µg	7(23%)	23(77%)	0(0%)
Sulphamethoxazole (SXT)	25µg	25(50%)	25(50%)	0(0%)

*Multiple antibiotic resistance (MAR) Index of *Edwardsiella tarda* strains (n = 30) isolates in this study*

The table below shows the number of resistant antibacterial from all *Edwardsiella tarda* isolates (n=30) and the multiple antibiotic resistance index of all the isolates. Multiple Antibiotic Resistance index is used to determine the

high risk of contamination with the use of antibiotics that pose a high risk to humans and also the prevalence of antibiotics in the treatment of *Edwardsiella tarda* bacteria. Multiple Antibiotic Resistance (MAR) index for each isolate was identified according to Singh *et al.* (2010), MAR index = Number of resistant antibiotics in a

single isolate (Number of antibiotics to which the bacterial isolates showed resistance) / Total Number of tested antibiotics. A Multiple Antibiotic Resistance index value of equal to or less than 0.2 was defined as those antibiotics that were seldom or never used for the animal in terms of treatment, whereas a MAR index value higher than 0.2 is considered that the fish has received high-risk exposure to those antibiotics. Therefore, it can be deduced from the table below that the fishes had received a high risk of exposure to antibiotics commonly used in aquaculture practices as all the isolates showed a MAR index ranging from 0.3 to 0.4. All isolates showed a MAR index of 0.4 except for F27 which showed a MAR index of 0.3 with a total of three antibiotics showing resistance to the isolate. The average MAR index for all the tested isolates was 0.397 (Table 3) (Figs. 1 to 3).

*Antimicrobial susceptibility profile of *Edwardsiella tarda* isolates (n = 30) isolated in this study also show their multiple antibiotic resistance index (MAR)*

The table below shows the susceptibility profile of *Edwardsiella tarda* isolates to ten commonly used antibiotics in aquaculture, the multiple antibiotics resistance index (MAR) was also determined from the number of antibiotics that were resistant to the isolate(s)/ total number of antibiotics tested. MAR index greater than 0.2 is considered that the fish had received high-risk exposure to those antibiotics. Therefore, 97% of *Edwardsiella tarda*

isolates showed an MAR index of 0.4 while only 3% of *Edwardsiella tarda* isolates showed a MAR of 0.3 indicating that the fish were highly exposed to the antibiotics that were tested on the isolates. The mean value of the MAR index showed a value of 0.397, *Edwardsiella tarda* isolates showed complete resistance to tetracycline, ampicillin and cefotaxime while showing a 90% resistance to chloramphenicol. *Edwardsiella tarda* was susceptible to Gentamycin, Enfloxacin and Norfloxacin, indicating that Gentamycin, Enfloxacin and Norfloxacin are effective in controlling *Edwardsiella tarda* infections (Table 4).

Table 3: Multiple Antibiotic Resistance (MAR) Index of *Edwardsiella tarda* strains (n = 30) isolates in present study.

Isolates	No resistant antibacterial	MAR index
F1	4	0.4
F2	4	0.4
F3	4	0.4
F4	4	0.4
F5	4	0.4
F6	4	0.4
F7	4	0.4
F8	4	0.4
F9	4	0.4
F10	4	0.4
F11	4	0.4
F12	4	0.4
F13	4	0.4
F14	4	0.4
F15	4	0.4
F16	4	0.4
F17	4	0.4
F18	4	0.4
F19	4	0.4
F20	4	0.4
F21	4	0.4
F22	4	0.4
F23	4	0.4
F24	4	0.4
F25	4	0.4
F26	4	0.4
F27	3	0.3
F28	4	0.4
F29	4	0.4
F30	4	0.4
Average		0.397

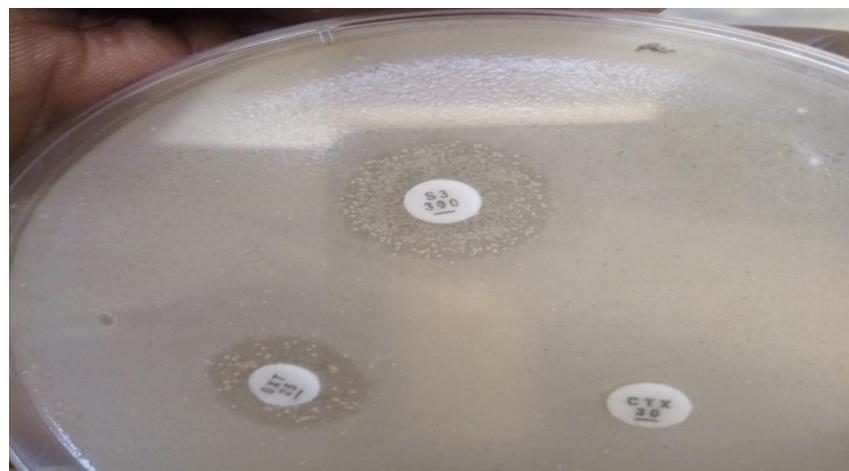


Figure 1: Antimicrobial susceptibility disk containing Sulphamethoxazole (SXT), Composulphonamides (S3) and Cefotaxime (CTX) Antibiotics containing a disk content of 25 μ g, 30 μ g and 30 μ g, respectively.

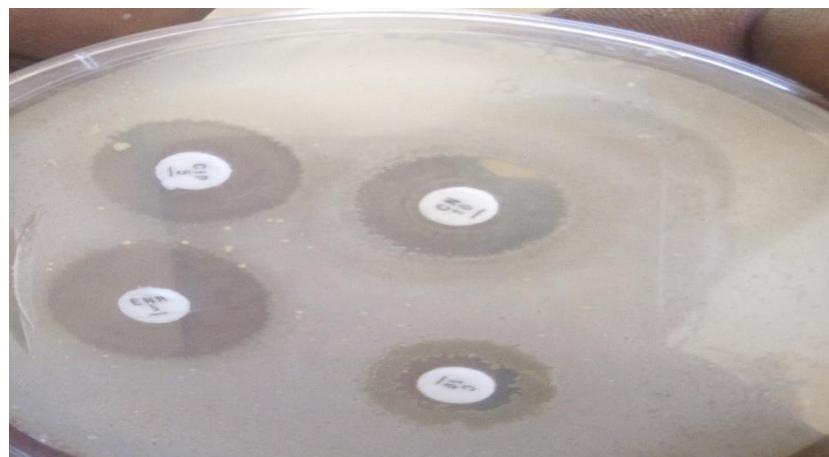


Figure 2: Antimicrobial susceptibility disk containing Enfloxacin (ENR), Gentamycin (CN), Chloramphenicol (C) and Ciprofloxacin (CIP) Antibiotics containing a disk content of 5 μ g, 10 μ g, 30 μ g and 5 μ g, respectively.

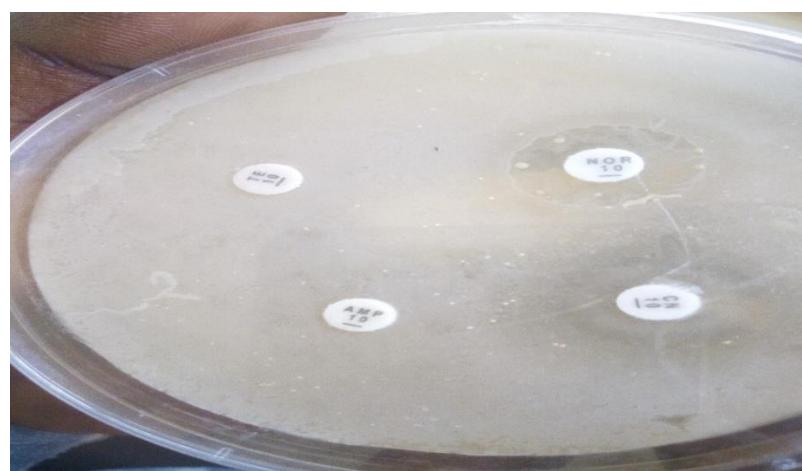


Figure 3: Antimicrobial susceptibility disk containing Tetracycline (TE), Ampillicin (AMP), and Norfloxacin (NOR) Antibiotics containing a disk content of 10 μ g, 10 μ g, and 10 μ g, respectively.

Table 4: Antimicrobial susceptibility profile of *Edwardsiella tarda* isolates (n = 30) isolated in this study also showing their multiple antibiotic resistance index (MAR).

	F1	F2	F3	F4	F5	F6	F7	F8	F9	F10	F11	F12	F13	F14	F15	F16	F17	F18	F19	F20	F21	F22	F23	F24	F25	F26	F27	F28	F29	F30
ENR	S	S	S	S	I	S	I	S	S	S	S	S	S	S	S	S	S	S	S	S	S	I	I	S	I	S	S	S	I	
CN	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S	S		
C	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	I	S	I	I	
CIP	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	I	R	R
TE	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
AMP	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
NOR	I	I	I	I	I	S	S	S	S	S	S	S	S	S	I	I	S	S	I	I	S	I	S	S	S	S	S	S	S	
CTX	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	R	
S3	I	I	I	I	I	I	I	I	I	I	I	I	I	I	S	S	I	I	S	I	I	I	I	I	I	I	I	S	S	
SXT	I	I	I	I	I	I	I	I	S	S	I	S	I	I	S	I	I	I	S	S	S	S	S	S	S	S	S	S		
MAR	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.4	0.3	0.4	0.4	0.4	0.4	

KEYS

R = Resistance

S = Susceptible or Sensitive

I = Intermediate

MAR = Multiple Antibiotic Resistance Index

F1: F30 = Isolates

ENR: Enfloxacin, CN: Gentamycin, C: Chloramphenicol, CIP: Ciprofloxacin, TE: Tetracycline, AMP: Ampicillin, NOR: Norfloxacin, CTX: Cefotaxime, S3: Composulphonamides, SXT: Sulphamethoxazole

Discussion

Antimicrobial agents can be used as a tool to maintain the health and disease prevention of cultured animals (Bischoff *et al.*, 2005). However, overuse of antimicrobial drugs has the potential to increase the occurrence of antibiotic resistance in harmful bacteria, lowering their susceptibility to antibiotics. Therefore, this study was carried out to reveal the antimicrobial resistance pattern and antimicrobial resistance prevalence of *Edwardsiella tarda*, an important bacterial disease in freshwater fish. *Edwardsiella tarda* is considered as one of the most important bacterial microorganisms that cause severe economic losses due to morbidity and mortality among various populations and age groups of fish in many countries. It is considered a dangerous septicemic pathogen with high economic losses and

highly encountered causes of diseases in stressed warm water aquaculture (Jun and Yin, 2006; Ibrahim *et al.*, 2011).

Antimicrobial sensitivity test in this study revealed that isolated *Edwardsiella tarda* was 100% resistant to Tetracycline, Ampilicin and Cefotaxime while 29 of the *Edwardsiella tarda* isolates showed 100 % resistance to Ciprofloxacin with a total resistance of 97%, Cefotaxime showed a resistance to *Edwardsiella tarda* isolates which was also agreed to (Nagy *et al.*, 2018) from their study justifying the case that cefotaxime is also a resistant antibiotic. This result indicates multiple antibiotic resistance and a total resistance to commonly used antibiotics in aquaculture such as Tetracycline, Ampilicin and Ciprofloxacin which are the frequently administered antibiotics used for treating

infections in humans and also in aquaculture. Wimalasena *et al.* (2018) reported that resistant isolates were at high frequency for the β -lactams antibiotics which are ampicillin and cefotaxime but at low frequency for the aminoglycosides (tetracycline) while Abd El Tawab *et al.* (2020) recorded a 75% resistance to ampicillin. Ogbonne *et al.* (2018) reported that *Edwardsiella tarda* showed a 100% resistance to Amoxycillin, which is under the same group of β -lactams antibiotics. *Edwardsiella tarda* isolates were also reported to be 100% resistant to chloramphenicol by Ogbonne *et al.* (2018), which was in contrast with this study, where chloramphenicol showed no resistance to *Edwardsiella tarda* but a 90% intermediate range to *Edwardsiella tarda* which is similar to the work of (Nagy *et al.*, 2018). Abd El Tawab *et al.* (2020) reported that Ciprofloxacin was 100% susceptible to *Edwardsiella tarda*, which was in agreement with (Ogbonne *et al.*, 2018). Whereas, the study here showed that Ciprofloxacin was 96.67% resistant to *Edwardsiella tarda*, with an indication that *E.tarda* is highly resistant to Ciprofloxacin. Tetracycline was found to be 100% resistant to *Edwardsiella tarda* in this study, while Abd El Tawab *et al.* (2020) reported 25% resistance of tetracycline to *Edwardsiella tarda* isolates in Egypt.

Sulphamethoxazole showed a 50% intermediate-range and 50% sensitivity range to *Edwardsiella tarda*, which was different from the study of (Nagy *et al.*, 2018), where the antibiotic was resistant

to the isolates. Abd El Tawab *et al.* (2020) reported that Sulphamethoxazole showed a 100% susceptible range to *Edwardsiella tarda* which was different from the result in this study where Sulphamethoxazole showed a 50% intermediate sensitivity range and 50% sensitivity range to *Edwardsiella tarda* indicating that the bacteria is developing resistance to the antibiotic over time due frequent and uncontrolled usage. The study carried out shows that Gentamycin, Enfloxacin and Norfloxacin were susceptible to *Edwardsiella tarda* isolates which were in close agreement with the result of Ogbonne *et al.* (2018) and Abd El Tawab *et al.* (2020). This therefore implies that Gentamycin, Enfloxacin and Norfloxacin are effective in controlling *Edwardsiella tarda* infections.

However, our results showed resistance to more than one class of antibiotic, and this is in agreement with the work of (Lee *et al.*, 2011), whose study reported multiple drug resistance by *Edwardsiella tarda* in freshwater fish, indicating that the farmed fishes within the Ilorin metropolis have high exposure to antibiotics.

Conclusion

The result of the present study shows the isolation of *Edwardsiella tarda* from commonly sold farmed fish in Ilorin Metropolis; the study also shows that *Edwardsiella tarda* is resistant to some commonly used antibiotics in aquaculture practices.

The study shows the prevalence of antimicrobial resistance in *Edwardsiella tarda* from farmed fishes in Ilorin Metropolis, indicating that farmed fishes in Ilorin Metropolis had a high risk of exposure to antibiotics.

References

Abd El Tawab A.A., El-Hofy, F.I., El-Gohary, M. and Sedek, A., 2020. Benha Veterinary Medical Journal antibiotic resistance genes of *Edwardsiella tarda* isolated from *Oreochromis niloticus* and *Clarias gariepinus*.
<https://bvmj.journals.ekb.eg/article>

Abu, G.O. and Wondikom, A.C., 2018. Isolation, Characterization and Antibiotic Resistance Profile Studies of Bacteria from an Excavated Pond in Port Harcourt Metropolis, Nigeria. *Journal of Applied Science and Environmental Management*, 22(8), 1177–1184.
<https://doi.org/10.4314/jasem.v22i8.6>

Bauer, A.W., Kirby, W.M., Sherris, J.C. and Truck, M., 1966. Antibiotic Susceptibility testing by a standardized disk method. *Journal Clinical Pathology*, 36, 493–496.
https://doi.org/10.1093/ajcp/45.4_ts.493

Bischoff, K.M., White, D.G., Hume, M.E., Poole, T.L. and Nisbet, D.J., 2005. The chloramphenicol resistance gene is disseminated on transferable plasmids that confer multiple-drug resistance in swine *Escherichia coli*. *FEMS Microbiology Letters*, 243, 285–291.

<https://doi.org/10.1016/j.femsle.2004.12.017>

CLSI, 2020. M100 Performance Standards for Antimicrobial Susceptibility Testing A CLSI supplement for global application. 30th Edition.
<https://www.nih.org.pk/wp-content/uploads/2021/02/CLSI-2020.pdf>

Ewing, W.H., McWhorter, A.C., Escobar, M.R. and Lubin, A.H., 1965. *Edwardsiella*, a new genus of Enterobacteriaceae based on a new species, *Edwardsiella tarda*. *International Journal System Evolution Microbiology*, 15, 33–38.
<https://doi.org/10.1099/00207713-15-1-33>

Ibrahem, M.D., Iman, B., Shaheed, H.A. and Korani, H., 2011. Assessment of the susceptibility of polyculture reared African Catfish and Nile tilapia to *Edwardsiella tarda* *Journal of American Science*, 7(3), 779–786.
<http://www.americanscience.org/>

Ibrahim, M., Ahmad, F., Yaqub, B., Ramzan, A., Imran, A., Afzaal, M., Mirza, S. A., Mazhar, I., Younus, M., Akram, Q., Ali Taseer, M. S., Ahmad, A. and Ahmed, S., 2020. Current trends of antimicrobials used in food animals and aquaculture. *Antibiotics and Antimicrobial Resistance Genes in the Environment*, 1, 39–69.
<https://doi.org/10.1016/b978-0-12-818882-8.00004-8>

Jun, Z. and Yin, L., 2006. Functional studies of type III and a novel

secretion system in *Edwardsiella tarda*. Ph.D. Thesis, National University of Singapore.

Jun, Z., 2006. Functional studies of a type III and a novel secretion system of *Edwardsiella tarda* (Doctoral dissertation).
<https://scholarbank.nus.edu.sg/handle/10635/15713>

Lee, S.W., Najiah, M., Chuah, T.S., Noor azhar, M.S., Wendy, W., Nadira, M. and Mohd, E.A.W., 2011. Antibiogram and Plasmid Profiling from *Edwardsiella tarda* Isolated from Freshwater Fish in East Coast Malaysia. *Journal of Sustainability Science and Management*, 6, 19-27.
<https://jssm.umt.edu.my/files/2012/01/3.June11.pdf>

Ma, F., Xu, S., Tang, Z., Li, Z. and Zhang, L., 2021. Use of antimicrobials in food animals and impact of transmission of antimicrobial resistance on humans. *Biosafety and Health*, 3(1), 32–38.
<https://doi.org/10.1016/j.bsheal.2020.09.004>

<https://doi.org/10.1016/j.bsheal.2020.09.004>

Manyi-Loh, C., Mamphweli, S., Meyer, E. and Okoh, A., 2018. Antibiotic Use in Agriculture and Its Consequential Resistance in Environmental Sources: Potential Public Health Implications. *Molecules*, 23, 795 – 843.
<https://doi.org/10.3390/molecules23040795>

Mohanty, B.R. and Sahoo, P.K., 2007. Edwardsiellosis in Fish: A Brief Review. *Journal of Bioscience*, 32, 1331-1344.
<https://doi.org/10.1007/s12038-007-0143-8>

Nagy, E., Fadel, A., Al-Moghny, F.A. and Ibrahim, M.S., 2018. Isolation, Identification and Pathogenicity Characterization of *Edwardsiella tarda* Isolated From *Oreochromis niloticus* Fish Farms in Kafr-Elshiekh, Egypt. *Alexandria Journal for Veterinary Sciences*, 57(1).
<https://doi.org/10.5455/ajvs.294237>

Nyandjou, Y.M.C., Yakubu, S.E., Abdullahi, I.O. and Machido, D.A. 2019. Multidrug Resistance Patterns and Multiple Antibiotic Resistance Index of *Salmonella* Species Isolated from Waste Dumps in Zaria Metropolis, Nigeria. *Journal of Applied Science and Environmental Management*, 23(1), 41–46.
<https://dx.doi.org/10.4314/jasem.v23i1.6>

Ogbonne, F.C., Ukazu, E.R. and Egbe, F.C., 2018. Antibiotics Resistance Pattern and Plasmid Profiling of *Edwardsiella tarda* Isolated from *Heterobranchus longifilis*. *Journal of Biosciences and Medicines*, 6, 95-105.
<https://doi.org/10.4236/jbm.2018.64008>

Osundiya, O., Oladele, R. and Oduyebi, O., 2013. Multiple Antibiotics Resistance (MAR) indices of *Pseudomonas* and *Klebsiella* species isolates in Lagos University Teaching Hospital. *African Journal of Clinical and Experimental Microbiology*, 14(3),

164–168.
<https://doi.org/10.4314/ajcem.v14i3.8>

Singh, A., Yadav, S., Singh, S. and Bharti, P., 2010. Prevalence of *Salmonella* in chicken eggs collected from poultry farms and marketing channels and their antimicrobial resistance. *Food Research International*, 43, 2027-2030. <https://doi.org/10.1016/j.foodres.2010.06.001>

Wimalasena, S.H.M.P., Pathirana, H.N.K.S., De Silva, B.C.J., Hossain, S., Sugaya, E., Nakai, T. and Heo, G.J., 2018. Antibiotic resistance and virulence-associated gene profiles of *Edwardsiella tarda* isolated from cultured fish in Japan. *Turkish Journal of Fisheries and Aquatic Sciences*, 19(2), 141-148. http://doi.org/10.4194/1303-2712-v19_2_06

Zhou, G., Shi, Q.-S., Huang, X.-M. and Xie, X.B., 2015. The Three Bacterial Lines of Defense against Antimicrobial Agents. *International Journal of Molecular Sciences*, 16(9), 21711–21733. <https://doi.org/10.3390/ijms1609217>