

ANTIBIOGRAM OF ESBL-PRODUCING ENTEROBACTERIACEAE ISOLATED FROM FISH COLLECTED FROM INTEGRATED POULTRY-FISH FARMS AT ILA ORANGUN, OSUN STATE, NIGERIA

^{*1}Adeluwoye-Ajayi, Olayemi Amos., ²Akinnibosun, Faith Iguodala.,
³Akinnibosun, Henry Adewale., ⁴Iyekowa, Osaro., ⁵Akinnibosun, Adeyemi Osarodion

¹Department of Science Laboratory Technology (Microbiology Option), Rufus Giwa Polytechnic, Owo, P.M.B 1019, Ondo State Nigeria.

²Department of Microbiology, Faculty of Life Sciences, University of Benin, P. M. B. 1154, Benin City, Nigeria.

³Department of Plant Biotechnology, Faculty of Life Sciences, University of Benin, P. M. B. 1154, Benin City, Nigeria.

⁴Department of Chemistry, Faculty of Physical Sciences, University of Benin, P. M. B. 1154, Benin City, Nigeria.

⁵School of Medicine, College of Medical Sciences, University of Benin, P. M. B. 1154, Benin City, Nigeria.

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Abstract: Antimicrobial resistance (AMR) is widely recognized as a silent pandemic, threatening decades of medical progress and complicating the treatment of infectious diseases worldwide. Among the myriad resistance determinants, extended-spectrum β -lactamases (ESBLs) have emerged as a major driver of therapeutic failure in clinical medicine. This study is aimed at determining the pattern of resistance of ESBL-producing enterobacteriaceae isolated from fish collected from integrated poultry-fish farms at Ila Orangun, Osun State, Nigeria. From November 2024 to April 2025, a total of 18 catfish were aseptically and randomly collected from six poultry-fish farms, processed and bacteriologically analysed using standard microbiological techniques. Isolated isolates were conventionally characterized, ESBL-production in them were confirmed via double-disc synergy test, and antibiogram of the confirmed ESBL+ isolates were carried out via Kirby-Bauer disc diffusion method. Out of 47 isolates recovered from the catfish, 2 (4.26%) were ESBL+. The ESBL+ isolates were totally resistance (100%) to the beta-lactam combination, the cepheems and the monobactam respectively tested in this study but totally susceptible (100%) to the carbapenem. They were totally (100%) multidrug-resistance (MDR) with 50% of them resisting 5 and 4 classes of antibiotics respectively, and showing 0.6 and 0.4 multiple antibiotic resistance index (MARI) respectively. The results confirm integrated agro-ecosystems as important reservoirs of MDR, ESBL-producing Enterobacterales. Urgent interventions in the farming practices, surveillance, and public education are needed to mitigate MDR, ESBL-producing enterobacteriaceae spread from farms produce to human through the food chain.

Keywords: Enterobacterales, Antimicrobial resistance, Extended-spectrum β -lactamases, Pandemic, Catfish.

1. INTRODUCTION

Fish is low in cholesterol and contains all essential amino acids and is estimated to provide roughly 60% of the world's protein requirements, with 60% of the developing world obtaining more than 30% of their animal protein from fish (Kuton *et al.*, 2021). In addition to being a good source of protein, fish is high in vitamins, unsaturated fatty acids, and essential minerals (Rajeshkumar & Li, 2018). The world's focus has shifted towards fish to address the deficiency of animal protein,

particularly in light of the increasing global population and depletion of natural resources (Du Thanh *et al.*, 2017). The high consumption of fish also stems from the fact that it is cheaper and generally accepted without any religious bias giving it an added benefit over other sources of proteins (Aliyu *et al.*, 2018). The Food and Agriculture Organization (FAO) estimates an increase in fish production in Nigeria by 2030 to about 18.2%, while export will increase by about 6.6% (FAO, 2018).

Cases of fish-borne disease caused by pathogenic microorganisms reported in several parts of the world heightened consumers fear regarding fish meat and fish products (Li *et al.*, 2019; Yanestria *et al.*, 2019). The use of organic manure also leads to the release of high concentration of opportunistic and pathogenic microorganisms into the ponds which pose a threat not only to fish health but the environment (Abdel-Wahed *et al.*, 2018). Also, these microorganisms in fish and fish ponds portend grave consequences for public health (Li *et al.*, 2019; Yanestria *et al.*, 2019).

Antimicrobial resistance (AMR) is widely recognized as a silent pandemic, threatening decades of medical progress and complicating the treatment of infectious diseases worldwide (Murray *et al.*, 2022). Both pathogenic and non-pathogenic bacteria can exhibit AMR. Multiple antibiotic resistance (MAR) is a worldwide problem caused by the abuse of antibiotics (Sharma *et al.*, 2015). Multidrug resistance (MDR) has increased all over the world and is considered a public health threat (Catalano *et al.*, 2022). The presence of MDR bacteria in fish can endanger public health because fish are thought to be potential carriers of foodborne bacterial diseases (Hassen *et al.* 2020). From investigations reported, the emergence of multidrug-resistant bacterial pathogens from different origins, increases the necessity of the proper use of antibiotics, routine applications of the antimicrobial susceptibility testing to detect the antibiotic of choice, and screening of emerging MDR strains (Abolghait *et al.*, 2020; Algammal *et al.*, 2020; Algammal *et al.*, 2022).

Among the myriad resistance determinants, extended-spectrum β -lactamases (ESBLs) have emerged as a major driver of therapeutic failure in clinical medicine as they hydrolyze a broad spectrum of β -lactams, including third-generation cephalosporins such as cefotaxime, ceftriaxone, and ceftazidime, leaving limited treatment options (Bush & Bradford, 2020), thus ESBL-producing bacteria can exhibit co-resistance to multiple classes of antibiotics (Tacão *et al.*, 2014). Infarct, the emergence of ESBL-producing *Enterobacteriaceae* became one of the crucial public health concerns, and subsequently fewer treatment options (Castanheira *et al.*, 2021).

Historically, ESBL-producing Enterobacterales were considered a nosocomial concern; however, in the past two decades, their epidemiology has shifted, with increasing prevalence in community, animal, and environmental settings (Bevan *et al.*, 2021). This reflects the interconnected nature of AMR as described in the One Health framework, which emphasizes the interdependence of human, animal, and environmental health (Robinson *et al.*, 2016). Food production systems, particularly integrated farms where poultry and fish intersect, are emerging as high-risk hotspots for resistance dissemination (Van Boeckel *et al.*, 2019). High organic loads and possible antibiotic residues from veterinary use in this system create selective pressures that favor resistant population (Gothwal & Shashidhar, 2020).

Although integrated poultry-fish farming systems are economically efficient but may inadvertently recycle resistant bacteria and ARGs across compartments, creating opportunities for horizontal gene transfer (HGT) and amplification of resistance determinants. The generation of extended-spectrum β -lactamase (ESBL) by *Enterobacteriaceae* can intensify this challenge (Iredell *et al.*, 2016).

There is insufficient data on the presence and spread of drug resistant ESBL-producing enterobacteriaceaea in integrated poultry-fish farms in Nigeria hence this study aimed at determining the pattern of resistance of ESBL-producing enterobacteriaceaea isolated from fish collected from integrated poultry-fish farms at Ila Orangun, Osun State, Nigeria so as to enhance antibiotic resistance control efforts.

2. MATERIALS AND METHODS

Description of Study Area

The study was conducted in commercialised integrated poultry-fish farms system located at Ila Orangun, Osun State, Nigeria. Ila Orangun, the headquarters of Ila Local Government Area in Osun State Nigeria, has an area of approximately 303km² and a population of 62,049 (2006 census). However, according to the information obtained from GeoNames geographical database, the population of Ila Orangun is 179, 192. The latitude and longitude coordinate of this town are 8.019116 and 4.901962 respectively. A common traditional profession of the indigenes of the town is palm-wine tapping. The people of Ila speak the distinctive dialect of the Yoruba language called Igbomina (or Igbonna).

Consideration of Ethical Issues

The study's purpose was explained to the integrated poultry-fish farmers, who consented to sampling on the condition that their identities not be published, hence the sampling farms are anonymous in this report. This study was performed in accordance with guidelines of the University of Benin research committee and the guidelines for the care and use of Agricultural Animals in research and teaching (Grandin *et al.*, 2020).

Samples Collection

Between November 2024 and November 2025, with the help of farms' manager, eighteen of five to six months old (adult) live apparently healthy African catfish (*Clarias gariepinus*) with approximately similar wet weight (1.5kg to 2.5kg) and length (29cm to 35cm) were randomly collected with the aid of cast net into bowls containing water from the fishponds of origin. Samples were collected once in two months (to give room for samples processing, bacteria isolation and identification) in triplicates using triangular method of sampling from the same integrated system on each of the six farms early in the morning (06:00am to 09:00am) for the period of sampling. All samples collected from each of the three integrated system per farm were well labelled and transported to the laboratory for immediate processing and bacteriological analysis.

Sample Preparation for Microbiological Study

The sample was prepared by making dilutions in distilled water.

Bacteriological Analysis

Bacterial isolation was done according to the method described by Cheesbrough (2016). The media used were prepared according to manufacturer's instruction. The glass wares such as Petri dishes, conical flasks, test tubes, beakers and bijou bottles were thoroughly washed and sterilized in a hot air oven at 160 °C for an hour. The inoculating loop was sterilized by flaming in the Bunsen burner until it turns red hot. Similarly, microbial load on the working surfaces were reduced by the application of disinfectant solution (70% ethanol).

Maintenance of Bacteria Stock Culture

Each of the isolates were maintained as stock cultures for further studies on nutrient agar (Oxoid, UK) slants in Bijou bottles. This was done by streaking the isolates on the agar slants, and incubating at 37 °C for 24 hrs. After incubation, the inoculated slants were stored in the refrigerator at ambient temperature and these stock cultures served as source of each isolate for further bacteriological studies.

Identification of Bacterial Isolates

Morphological Characterization of Isolates

A 24-hour old pure culture of the isolates was characterized and the different morphologies were recorded.

Gram-Staining and Microscopic Examination

This was carried out according to the method described by (Becerra *et al.*, 2016)

Biochemical Characterization of the Isolates

The biochemical tests such as motility test, catalase test, citrate utilization test, indole test, methyl red test, voges-proskauer test, and sugar fermentation (glucose, lactose and maltose) test were carried out according to Cheesbrough (2016).

Screening for Potential Extended Spectrum Beta-Lactamase Producing Enterobacteriaceae

All the isolates were assessed for the production of extended spectrum beta-lactamase. Mueller-Hinton agar plates were inoculated with homogeneous inoculum of the test isolates and separate antibiotic discs that contain ceftazidime (30 µg) and cefotaxime (30 µg) respectively were positioned aseptically on the agar surface at a distance of 30 mm apart. The plates were placed on workbench for about half-hour to allow the antibiotics to diffuse into the agar, and then they were incubated at 37 °C for 24 hrs. Thereafter, meter rule was used to measure the zones of inhibition. A reduced susceptibility/resistance to either of the two antibiotics used for the screening was taken as suspected case of extended spectrum beta-lactamase production (CLSI, 2024).

Recognition of Extended Spectrum Beta-Lactamase Producing Enterobacteriaceae

An 18 hrs old culture of isolates that shows resistance to one or more of the third generation cephalosporins were inoculated into test tubes which contain sterile normal saline, then the turbidity of the content was adjusted to 0.5 MacFarland standards. The bacterial suspension was inoculated on prepared Mueller Hinton agar plates by uniformly swabbing the entire surface of the agar plates. The double disc synergy test was done using discs of augmentin alongside the discs of ceftazidime (30 µg) and cefotaxime (30 µg); which were placed at 20 mm equidistance from augmentin disc. Then, the plates were incubated for 24 hrs at 37 °C, the plates were observed and the organisms was considered to be producing extended spectrum beta-lactamase when the zone of inhibition is ≤5 mm in diameter of either cefotaxime or ceftazidime in combination with augmentin against its zone when tested alone (CLSI, 2024)

Antibiogram of the ESBL-producing Bacteria

This was done using the standard Kirby-Bauer disc diffusion (Jayabarath, 2015). The ESBL-producing bacteria inoculum was prepared by separately suspending the freshly grown bacteria in 5 ml sterile nutrient broth and its turbidity adjusted to 0.5 McFarland standards. The antimicrobial susceptibility testing was performed separately on Mueller-Hinton agar using the following antibiotics: beta-lactam combination agent (augmentin 20/10µg), cephem (cefotaxime 30µg, ceftazidime 30µg, cefixime 5µg), carbapenem (imipenem 10µg), aminoglycosides (gentamicin 10µg), fluoroquinolone (ciprofloxacin 5µg, ofloxacin, 5µg), monobactam (aztreonam 30µg) and nitrofurans (nitrofurantoin 300µg). The plates were incubated aerobically at 37 °C for 24 h. The zones of inhibition were measured with a metre rule and the results were recorded and interpreted according to the Clinical and Laboratory Standards Institute (CLSI) guidelines (2024). Isolates showing resistance to at least three different classes of antibiotics were considered as multidrug resistant strains (Magiorakos *et al.*, 2012). *E. coli* ATCC 25922 was used as a quality control organism

Multiple Antibiotic Resistance Index of Isolates

As described by Ekwealor *et al.* (2016), the calculation of the Multiple Antibiotic Resistance Index (MARI) involved employing the provided equation below. This index was derived by assessing whether a zone of inhibition was present or absent. The MARI value was determined by dividing the total count of antibiotics to which an isolate demonstrated resistance by the total count of antibiotics administered to each strain.

$$MARI = \frac{\text{Number of antibiotics strain resistant}}{\text{Number of antibiotics strain subjected}}$$

3. RESULTS

Morphological, Macroscopic, Cultural and Staining Characteristics of the Bacteria Isolated from the Catfish Collected from the Integrated Poultry-Fish Farms

Table 1 below showed the morphological, macroscopic, cultural and staining characteristics of the bacteria isolated from catfish collected from the six integrated poultry-fish farms at Ila Orangun, Osun State, Nigeria. Inferences drawn from the bacteria characteristics are as depicted in the table.

Biochemical Characteristics of the Bacteria Isolated from the Samples Collected from the Integrated Poultry-Fish Farms

Table 2 below showed the biochemical characteristics of the bacteria isolated from catfish collected from the six integrated poultry-fish farms at Ila Orangun, Osun State, Nigeria. Inferences drawn from the biochemical characteristics were used in getting the identity of the isolates as depicted in the table.

Preliminary Screening for ESBL Producing Isolates from Catfish Samples from the Integrated Poultry-fish Farms

The preliminary screening for ESBL production by *E. coli*, *K. pneumonia*, *K. aerogenes*, *Salmonella sp.*, *Enterobacter sp.* and *Proteus sp.* isolated from catfish is shown in Table 3 where it was revealed that 7 (41.18%), 1 (8.33%), 1 (33.33%), 1 (25.0%), 1 (20.0%) and 1 (16.67%) of the respective isolates were suspected to be ESBL producers since they showed resistance to cefotaxime.

Confirmatory Screening for Suspected ESBL-producing Isolates from Catfish Collected from Poultry-fish Pond Integrated Farms

The confirmatory screening for ESBL production by isolates from catfish indicated that 2 (16.67) out of the 12 suspected ESBL positive isolates from catfish were confirmed to be ESBL positive. Furthermore, out of all the bacteria isolated only *E. coli* produced ESBL which was just 2 (28.57) of the 7 suspected *E. coli* (Table 4).

Prevalence of ESBL-Producing Enterobacterales in the Catfish from the Integrated Poultry-Fish Farms

Table 5 below showed that out of the 47 enterobacterales recovered from the fish sampled, only *E. coli* produced ESBL and its prevalence in relation to the total number of isolates isolated was just 4.26%

Antibiogram of the ESBL-Producing *E. coli* (n = 2) Isolated from Catfish Collected from Integrated Poultry-Fish Farms

Table 6 showed the antibiogram of the only ESBL-producing enterobacteriaceae (i.e. *E. coli*) isolated from catfish collected from the farms to the tested antibiotics. The ESBL-producing *E. coli* isolates from the sample were totally resistance (100%) to the beta-lactam combination (augmentin), the cepheims (cefotaxime, ceftazidime and cefixime) and the monobactam (aztreonam) respectively tested in this study but totally susceptible (100%) to the carbapenem (imipenem).

Antibiotype of ESBL Producing *E. coli* (n = 2) Isolated from Catfish Collected from

Integrated Poultry-Fish Farms

The antibiotypes of the ESBL producing *E. coli* showed that 1 (50%) of the two isolates that produced ESBL resisted five classes of antibiotics (CIP-CAZ-NIT-GEN-OFL-AUG) and four classes of antibiotics (CIP-CAZ-GEN-AUG) respectively, and showing 0.6 and 0.4 multiple antibiotic resistance index (MARI) respectively. The ESBL-producers were totally (100%) multidrug-resistant (Table 7).

Table 1: Morphological, Macroscopic, cultural and Staining Characteristics of the Bacteria Isolates from the Samples Collected from the Integrated Poultry-fish farms

Colony characteristics on MacConkey Agar	Morphology (Staining characters)	Probable organism
Smooth, circular, flat and non-mucoid, colonies that are pink to reddish-pink due to lactose fermentation	Gram-negative (pink/red colour), small (1.0–2.0 µm long and 0.5 µm wide), rod-shaped, arranged in single or paired short, motile, Non-sporing, some are capsulated	<i>E. coli</i>
Smooth, large (often 4-6 mm), round, convex (dome-shaped), and shiny colonies that are pink to dark pink/red due to lactose fermentation, appearing mucoid slimy, glistening, and sticky because of their prominent polysaccharide capsule.	Gram-negative (pink/red colour), small rod-shaped, arranged in single or paired short, Non-motile, Non-spore-forming, capsulated	<i>Klebsiella</i> sp.
small to medium (2-3 mm), convex (dome-shaped), smooth, and can have slightly serrated or entire margins, colorless or pale, translucent colonies because they are non-lactose fermenters	Gram-negative (pink/red colour), small (2-5 µm long, 0.8-1.5 µm wide) rod-shaped, arranged in single or paired short, motile, Non-spore-forming, capsulated	<i>Salmonella</i> sp.
Pink to red/rose-red colonies due to lactose fermentation (Lac+) and produce a capsule, making the colonies appear moist, mucoid (slimy/sticky and sticky, often surrounded by a hazy zone.	Gram-negative (pink/red colour), motile, small (1–5 µm long) straight rod-shaped bacteria (bacilli) appearing as singly, dispersed or scattered, facultative anaerobic, and non-spore forming, capsulated bacteria.	<i>Enterobacter</i> sp.
Translucent, colorless, non-lactose fermenting colonies that often exhibit a characteristic swarming (forming a thin, filmy layer) growth pattern, spreading across the plate in concentric rings with a fishy or ammonia-like smell, though they may appear as smaller, smooth, pale or grayish-white colonies if motility is inhibited.	Gram-negative (pink/red colour), motile, non-spore-forming, non-encapsulated, rod-shaped (bacilli) bacteria that are sometimes pleomorphic	<i>Proteus</i> sp.

Table 2: Biochemical Characteristics of the Bacteria Isolated from the Samples Collected from the Integrated Poultry-fish Farms

MOT	CAT	CIT	IND	MR	VP	GLU	LAC	MAL	ISOLATES IDENTITY
+	+	-	+	+	-	+	+	+	<i>Escherichia coli</i>
-	+	+	-	-	+	+	+	+	<i>Klebsiella pneumoniae</i>
+	+	+	-	-	+	+	+	+	<i>Klebsiella aerogenes</i>
+	+	+	-	+	-	+	-	+	<i>Salmonella sp.</i>
+	+	+	-	-	+	+	+	+	<i>Enterobacter sp.</i>
+	+	+	-	+	-	-	-	+	<i>Proteus sp.</i>

KEY: MOT = Motility, CAT = Catalase, CIT = Citrate, IND = Indole, MR = Methyl Red, VP = Voges-Proskauer, GLU = Glucose, LAC = Lactose, MAL = Maltose, - = negative, + = positive, +/- = variable

Table 3: Preliminary Screening for ESBL-producing Isolates from Catfish Samples of Integrated Poultry-fish Farms

Name of isolate	No. of isolate	Antibiotics used	No. (%) resistant	No. (%) sensitive
<i>E. coli</i>	17 (36.2)	Cefotaxime	7 (41.18)	10 (58.82)
		Ceftazidime	6 (35.29)	11 (64.71)
<i>K. pneumoniae</i>	12 (25.5)	Cefotaxime	1 (8.33)	11 (91.67)
		Ceftazidime	0 (0)	12 (100.0)
<i>K. aerogenes</i>	3 (6.4)	Cefotaxime	1 (33.33)	2 (66.67)
		Ceftazidime	0 (0)	3 (100.0)
<i>Salmonella sp.</i>	4 (8.51)	Cefotaxime	1 (25.0)	3 (75.0)
		Ceftazidime	0 (0)	4 (100.0)
<i>Enterobacter sp.</i>	5 (10.6)	Cefotaxime	1 (20.0)	4 (80.0)
		Ceftazidime	0 (0)	5 (100)
<i>Proteus sp.</i>	6 (12.8)	Cefotaxime	1 (16.67)	5 (83.33)
		Ceftazidime	0 (0)	6 (100.0)
TOTAL	47			

KEY: No. = Number, % = Percentage

Table 4: Confirmatory Screening for Suspected ESBL-producing Isolates from Catfish Samples collected from Poultry-fish Pond Integrated Farms

Name of Isolate Screened	No. of Isolates Screened	No. of ESBL + (%) Isolates	No. of ESBL - (%) Isolates
<i>E. coli</i>	7	2 (28.57)	5 (71.43)
<i>K. pneumoniae</i>	1	0 (0)	1 (100.0)
<i>K. aerogenes</i>	1	0 (0)	1 (100.0)
<i>Salmonella sp.</i>	1	0 (0)	1 (100.0)
<i>Enterobacter sp.</i>	1	0 (0)	1 (100.0)
<i>Proteus sp.</i>	1	0 (0)	1 (100.0)
TOTAL	12	2 (16.67)	10 (83.33)

KEY: No. = Number, % = Percentage

Table 5: Occurrence of ESBL Producers among the Bacteria Isolated from the Fish Samples Collected from Poultry-fish Pond Integrated Farms

Isolate	Isolate No	No. (%) of Probable ESBL Producers	No. (%) of Confirmed ESBL Producers	No. (%) of Non ESBL Producers
<i>E. coli</i>	17	7 (41.18)	2 (11.76)	15 (88.24)
<i>K. pneumoniae</i>	12	1 (8.33)	0 (0)	12 (100.0)
<i>K. aerogenes</i>	3	1 (33.33)	0 (0)	3 (100.0)
<i>Salmonella</i> sp.	4	1 (25)	0 (0)	4 (100.0)
<i>Enterobacter</i> sp	5	1 (20)	0 (0)	5 (100.0)
<i>Proteus</i> sp.	6	1 (16.67)	0 (0)	6 (100.0)
Total	47	12 (25.53)	2 (4.26)	45 (95.74)

KEY: No. =Number; % = percentage

Table 6: Antibigram of ESBL-producing *E. coli* (n = 2) Isolated from Catfish Collected from Integrated Poultry-fish Farms

Family of Antibiotics Tested	Name of Antibiotics Tested	Antibiotics Disc Code	Antibiotics Disc Concentrations (µg)	Interpretive Categories and Zone Diameter Breakpoint, Nearest Whole Number (mm)			
				S	R	S No. (%)	R No. (%)
Beta-lactam combination	Augmentin	AUG	30	≥ 18	≤ 13	0 (0)	2 (100)
Cephems	Cefotaxime	CTX	30	≥ 26	≤ 22	0 (0)	2 (100)
	Ceftazidime	CAZ	30	≥ 18	≤ 14	0 (0)	2 (100)
	Cefixime	CXM	5	≥ 16	≤ 20	0 (0)	2 (100)
	Imipenem	IMP	10	≥ 19	≤ 15	2 (100)	0 (0)
Aminoglycosides	Gentamicin	GEN	10	≥ 15	≤ 12	1 (50)	1 (50)
Fluoroquinolones	Ciprofloxacin	CIP	5	≥ 21	≤ 15	1 (50)	1 (50)
	Ofloxacin	OFL	5	≥ 16	≤ 12	1 (50)	1 (50)
Monobactam	Aztreonam	AZT	30	≥ 22	≤ 15	0 (0)	2 (100)
Nitrofurantoin	Nitrofurantoin	NIT	300	≥ 17	≤ 14	1 (50)	1 (50)

KEY: n/No. = Number, % = percentage, S = Susceptible, R = Resistant

Table 7: Antibiotypes of ESBL-producing *E. coli* (n = 2) Isolated from Catfish Collected from Integrated Poultry-fish Farms

Pattern No.	Antibiotypes	No. of Antibiotics (Classes)	No. of Antibiotics tested	No. (%) ESBL+ <i>E. coli</i>	MAR Index
1	NIT-IMP-CIP-CAZ-GEN-AUG-AZT	7 (7)	10	0 (0)	0
2	CIP-IMP-AZT-GEN-CRX-AUG	6 (6)	10	0 (0)	0
3	CIP-CAZ-NIT-GEN-OFL-AUG	6 (5)	10	1 (50)	0.6
4	NIT-CIP-CXM-GEN-AUG	5 (5)	10	0 (0)	0
5	CIP-CRX-GEN-OFL-AUG	5 (4)	10	0 (0)	0
6	IMP-CXM-GEN-OFL	4 (4)	10	0 (0)	0
7	NIT-CXM-IMP-AUG	4 (4)	10	0 (0)	0

8	CIP-CAZ-GEN-AUG	4 (4)	10	1 (50)	0.4
9	CIP-CXM-OFL-AUG	4 (4)	10	0 (0)	0
10	CIP-CAZ-GEN-AUG	4 (4)	10	0 (0)	0
11	CAZ-OFL-AUG	3 (3)	10	0 (0)	0
12	CXM-GEN-AUG	3 (3)	10	0 (0)	0
13	IMP-CAZ-CTX	3 (2)	10	0 (0)	0
Total MDR of ESBL+ Isolates No. (%)				2 (100)	

KEY: n/No. = Number, % = percentage

4. DISCUSSION

Antibiotic resistance has been causing serious problems to human medicine and also to animal, livestock, and veterinarians (Lawson, 2008). The heavy and off-label use of antibiotics has been reported to be a risk factor for the development and spread of beta-lactamase producing organisms (Chishimba *et al.*, 2016; Economou and Gousia, 2015). Therefore, the present study focus on investigating the pattern of resistance of ESBL-producing enterobacteriaceae (ESBL-PEN) isolated from fish collected from integrated poultry-fish farms at Ila Orangun, Osun State, Nigeria.

The morphological, macroscopic, cultural and staining characteristics of the bacteria isolated from the fishponds water in this present study were in agreement with previous researchers report (Akinribosun *et al.*, 2024; Osuntokun *et al.*, 2020; Okpalaji *et al.*, 2025; Nwabueze & Nwabueze, 2011; Ali, 2025; Bello *et al.*, 2019; Agwaranze *et al.*, 2025; Kanu *et al.*, 2024; Ajayi, 2012; Fakorede *et al.*, 2020).

Also, the biochemical characteristics of the members of enterobacteriaceae isolated from the fishponds water in this present study were in agreement with previous researchers report (Akinribosun *et al.*, 2024, Osuntokun *et al.*, 2020; Okpalaji *et al.*, 2025; Nwabueze & Nwabueze, 2011; Ali, 2025; Bello *et al.*, 2019; Agwaranze *et al.*, 2025; Kanu *et al.*, 2024; Ajayi, 2012; Fakorede *et al.*, 2020).

The results of this study indicating the presence of ESBL positive (11.76%) *E. coli* is in agreement with the earlier reports of Elhadi (2016) in Saudi Arabia, Hinthong *et al.* (2024) in Thailand and Dayie *et al.* (2024) in Ghana each of whom respectively reported the presence of ESBL-producing *E. coli* (27.2 %, 5.7 %, and 7.4 % respectively) in fish samples. However, the 11.76% ESBL producing *E. coli* (ESBL-PEC) observed in this study is lower than 21.2% but higher than 5.7% and 7.4% respectively reported by same authors. The reason for this disparity could be due the number of isolates studied, the species of fish studied and the level of animal exposure to antibiotics.

The overall ESBL-producers (4.26%) in this present study is in agreement with the 1.3% reported by Dayie *et al.* (2024) in Ghana. However, the 4.26% ESBL producers observed in this study is a little bit higher than the 1.3% reported by same author. The reason for this disparity could be due to differences in the method of screening for ESBL producers, due the number of isolates studied, the species of fish studied and the level of animal exposure to antibiotics. In addition, none of the other isolates in this present study produced ESBL, a report which negate the report of Singh *et al.* (2017) and Dayie *et al.* (2024) both of whom reported the production of ESBL by the other bacteria species. The reason for this disparity could be due the number of isolates studied, the species of fish studied and the level of animal exposure to antibiotics.

The presence of ESBL-PEC in catfish from the farms is of public health importance as it point toward a great health risk for the consumers in the area of study. The contamination of fish meant for public consumption raises a potential risk of infection among the consumers as well as the handlers of these fish with ESBL producing *E. coli* particularly when these fish are eaten without adequate cooking. It is important to note that catfish are used in making pepper soups in the studied area, a recipe which undergo minimal heat processing which are incapable of neutralizing the microbial contaminants on the catfish. On the other hand, the type of resistance caused by ESBLs usually enables the organism to be resistant to other forms of antibiotics, and this in effect makes antibiotic therapy ineffective (Reist *et al.*, 2013). Deaths due to ESBL-producing *E. coli* infections are reported to be three-times greater than those caused by *E. coli* that are incapable of producing ESBLs (Tekiner & Özpınar, 2016; Melzer & Petersen, 2007).

Antibiotic resistance presents a significant public health concern, as antibiotic-resistant bacteria can cause severe illness or even death, and can be challenging to treat (Blaak *et al.*, 2015; García-Vello *et al.*, 2020). This present study revealed

different degree of resistance by the only isolated ESBL-producers (i.e. *E. coli*) among the isolates to the tested commercially available antibiotics. Out of all the tested antibiotics, the isolated ESBL-PEC showed total resistance (100%) to the beta-lactam combination (augmentin), the cepheims (cefotaxime, ceftazidime and cefixime) and the monobactam (aztreonam) respectively tested in this study but totally susceptible (100%) to the carbapenem (imipenem).

The total resistance (100%) to cefotaxime and ceftazidime respectively observed in this present study support the report of Singh *et al.* (2017) where 97.63% and 91.12% of the ESBL-producing isolates were respectively resistance to these antibiotics. The total resistance (100%) of the ESBL-PEC to ceftazidime observed in this present study is parallel to the same value of resistance to same antibiotic earlier reported by Chancelle *et al.* (2023). Also, the total resistance (100%) of the ESBL-PEC to cefotaxime observed in this present study is parallel to the same value of resistance to same antibiotic earlier reported by Chancelle *et al.* (2023). However, the total resistance (100%) of the ESBL-PEC to cefotaxime is far higher than the values earlier reported in previous research studies (Dayie *et al.*, 2024; Adinortey *et al.*, 2020; Chibuike *et al.*, 2021; Agbeko *et al.*, 2022).

The total resistance (100%) shown by the ESBL-PEC in this present study to aztreonam and augmentin respectively is far higher than the 65.08% and 38.46% resistance respectively shown by isolated bacteria to these antibiotics as reported by Singh *et al.* (2017). The total resistance (100%) shown by the ESBL-PEC in this present study to aztreonam support the report of Chibuike *et al.* (2021) and Chancelle *et al.* (2023) who reported 100% resistance and 89.5% resistance respectively to same antibiotic. Also, the total resistance (100%) shown by the ESBL-PEC in this present study to augmentin is in conformity with the same value of resistance to same antibiotic reported by Chibuike *et al.* (2021).

The observed 50% resistance to gentamicin in this study is far higher than the values earlier reported in previous studies (Hinthong *et al.*, 2024; Dayie *et al.* 2024; Singh *et al.*, 2017; Agbeko *et al.*, 2022; Adinortey *et al.*, 2020; Chancelle *et al.*, 2023). The 50% resistance shown by the isolate to ofloxacin in this present study is similar but lower than the 57% and the 94.7% resistance shown by same organism to same antibiotic reported by Adinortey *et al.* (2020) and Chancelle *et al.* (2023) respectively. Also, the 50% resistance shown by the isolate in the present study to ciprofloxacin is similar but a little bit lower than the 63.2% resistance shown by same organism to same antibiotic reported in earlier study by Chancelle *et al.* (2023). However, the observed 50% resistance to ciprofloxacin in this present study by the isolate negate the 0% resistance shown by same organism to same antibiotic reported by Mwanza *et al.* (2021).

The none resistance (0%) shown by the ESBL-PEC in this present study to the carbapenem (imipenem) is parallel to the value of resistance to same antibiotics earlier reported in previous research (Dayie *et al.*, 2024; Chibuike *et al.*, 2021). However, the zero resistance shown by the ESBL-PEC to the tested carbapenem (imipenem) in this present study is lower than the 10.65% and 18.2% resistance respectively reported by Singh *et al.* (2017) and Chancelle *et al.* (2023). The reason for the disparity in the level of antibiotics activity observed in the present study in relation to previous studies could be due the number of isolates studied and the level of the fish exposure to antibiotics. The reasonably high level of activity of the imipenem against the ESBL-PEC isolates in this study support the view of carbapenems being the standard drug for the management of infections occasioned by some Gram negative bacteria including those provoked by extended-spectrum- β -lactamases (Dahiya *et al.*, 2015).

All the ESBL-PEC in this study were multidrug-resistant with 50% of them resisting five (CIP-CAZ-NIT-GEN-OFL-AUG) and four (CIP-CAZ-GEN-AUG) classes of antibiotics respectively supporting the report of Kimera *et al.* (2021) whose studied showed eight isolates resisting eight classes of antimicrobials; and the report of Homeier-Bachmann *et al.* (2021) where all the 25 ESBL-producing Enterobacteriaceae were reported as multidrug-resistant.

The ESBL-PEC in this study showed 0.6 and 0.4 Multiple Antibiotic Resistance Index (MARI) respectively. According to Thenmozhi *et al.* (2014), Multiple Antibiotic Resistance Index (MARI) determines the level of an isolate's resistance to the antibiotics tested, and a value of ≥ 0.2 indicates a high risk of contamination from sources with high antibiotic use hence it could be stated that catfish from the source studied is of high risk of antibiotics contamination and is not safe for human consumption.

5. CONCLUSION

This study showed the presence of multidrug resistant (MDR), ESBL-PEC in the catfish gotten from the farms. Highest resistance (100%) shown by this isolate was to augmentin, cefotaxime, ceftazidime, cefixime and aztreonam respectively while the least resistance (0%) was to imipenem.

6. RECOMMENDATION

Urgent interventions in the farming practices, surveillance, and public education are needed to mitigate MDR, ESBL-producing enterobacteriaceae spread from catfish to human through the food chain.

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