



## Original Research Article

# Prevalence of Tetracycline Resistant *Aeromonas Hydrophila* Isolated From diarrheic Patients in Hilla City, Iraq

Lubna Abdulmutaleb Al-Shalah<sup>\*</sup> Alaa Hani Al-Charrakh College of Medicine, University of Babylon, Hilla, IRAQ

\*E-mail:lubnamsa77@gmail.com

Accepted 13 October, 2015

#### **Abstract**

The aim of this study was to investigate the prevalence of tetracycline resistant *Aeromonashydrophila* isolated from clinical sources in Hilla city, Iraq. A total of 822 samples were collected from fecal specimens from patients. Samples were collected from those who suffering from diarrhea. The period of the research was from October 2013 to February 2014 at public health lab, Hilla city. Results of this study revealed that out of 822 fecal samples, 13 isolates (1.58%) were belonged to *Aeromonas spp*. However, other bacterial isolates belonged to other genera similar to *Aeromonas* were also recovered. Out of 13 *Aeromonas spp*., eight *A. hydrophila* isolates (61.53%) were obtained, while the other isolates were distributed as: four isolates of *A. salmoncidia* (30.76%), and one of *A. sobria*. Isolation and detection of *A. salmoncidia* species was first recorded in Iraq. The susceptibly of *A. hydrophila* (NO. 8) to several antibiotics was determined using disk diffusion test (DDT). Results showed that 5 isolates (62.5%) were Multi-drug resistant (MDR) and three isolates (S2, S3, and S5,) were sensitive to most of antibiotic classes tested. The MIC of *A. hydrophila* to tetracycline was also detected using ager dilution mothed according to CLSI guidelines. Results found that only 2 isolates (25%) were resistant to tetracycline. The MIC of these isolates ranged from 0.25-16 µg/ml. This result confirms tetracycline resistance by these isolates when tested using DDT.

<u>Key words:</u> *Aeromonashydrophila ,Aeromonassalomonacidia* , Tetracyclineresistance , diarrheic patients , Iraq.

#### الخلاصة

تهدف هذه الدراسة الى الكشف عن مدى انتشار بكتريا Aeromonashydrophila المقاومة للتتراسايكلين والمعزولة من عينات الخروج للمرضى لمختبر الصحة العامة في مدينة الحلة. تم جمع ٢٢٢ عينة خروج في مختبر الصحة العامة للفترة من تشرين الاول ٢٠١٣ الى شباط ٢٠١٤ وتم تشخيصها من خلال اجراء الاختبارات الزرعيةوالبايوكيميائية وتأكيد النتائج باستخدام نظام ٧٧١٤ ، حيث اظهرت النتائج انه من مجموع انتخيصها من خلال اجراء الاختبارات الزرعيةوالبايوكيميائية وتأكيد النتائج باستخدام نظام ٧٧٤٤ ، حيث اظهرت النتائج انه من مجموع معنه خروج ، ١٣ عينه فقط كانت عائدة لبكتريا . *Aeromonasspp (٥*٠١.%) ، على الرغم من انه النسب الاخرى تعود لأنواع بكتيرية الخرى قريبة بصفاتها من بكتريا الاختبارات الزرعيةوالبايوكيميائية وتأكيد النتائج باستخدام نظام البلمرة المتعدد PCR عينه فقط كانت عائدة لبكتريا . *Aeromonasspp (٥*٠.٥.%) ، على الرغم من انه النسب الاخرى تعود لأنواع بكتيرية الخرى قريبية بصفاتها من بكتريا الاختبارات الزرعيةوالبايوكيميائية وتأكيد والتشخيص باستخدام نظام البلمرة المتعدد PCR عينه خروج ، ١٣ عينه فقط كانت عائدة لبكتريا . *Aeromonasspp (٥*.٥.%) ، على الرغم من انه النسب الاخرى تعود لأنواع بكتيرية الإيروموناس (٢٠٢٣) عور قريبة بصفاتها من بكتريا المحقمات المظهرية والزرعيةوالبايوكيمياوية فقد وجد ان العزلات تعود للأنواع ضمن جادئ خاص ببكتريا الايروموناس (٢٠٢٠%) عود للأنواع ضما بلايروموناس : عنود قريبة بصفاتها من بكتريا الامن المظهرية والزرعيةوالبايوكيمياوية فقد وجد ان العزلات تعود للأنواع ضمن حس الايروموناس : عنور موناس (٢٠٢٠%) تعود للنوع المنوموناس : معنه معنه الايروموناس (٢٠٢٠%) تعود للنوع المحضوم بكتريا محمومهما معنه من الايروموناس (٢٠٢٠%) تعود للنوع المصفات المظهرية والزرعيةوالبايوكيمياوية فقد وجرم مان بكتروا الايروموناس (٢٠٢٠%) عود المروموناس (٢٠٢%) معامل موليان عاربايوكيمياوية فود الى الايوموناس (٢٠٢٠%) تعود للنوع المصادات الحياتية استخدام طريقة انتشار الاقراص في الاكار. اظهرت النائع مان ما ما لايرام ما ما لايرام في الاكار. اظهرت النائع مان ما ما ما لايرام في الكرام ما ما لاكار الفيرام في الاكار. اظهرت النائع ما ما ما ما ما لاينا ما ما ما مالمايوين الولى ما ما ما ما ما ما ما لاينا ما ما لايرالمايكيين والتي تماومة (٢٠٦٠%) لكثر من ثلائة أصناف من المضادات ا

الكلمات المفتاحية :ايروموناسهايدروفيلا، ايروموناسسالمونسيديا، مقاومة التتراسايكلين، الاسهال، العراق.

### Introduction

embers of the genus *Aeromonas*are facultatively Lanaerobic, non- spore forming, rod shaped oxidase positive, gram negative bacteria motile by polar flagellum, mesophilic and facultative anaerobic bacteria of family Aeromonadaceae whose natural habitat is in the aquatic environment. Some species are pathogenic for animals and humans. Aeromonasspecies are widely distributed in the aquatic environment, including raw and processed drinking water, and have been frequently isolated from various food products such as fish and shellfish, raw meat, vegetables, and raw milk Additionally, in recent years aeromonads have been implicated as causative agents of human disease, ranging from gastroenteritis to wound infections [1-2].

The *Aeromonas* comprises genus important human pathogens causing primary and secondary septicemia in immunocompromised persons. serious wound infections in healthy individuals and in patients undergoing medicinal leech therapy, and a number of less well described illnesses such as peritonitis. meningitis, and infections of the eye, joints, and bones. Gastroenteritis, the most common clinical manifestation, remains controversial [3].

Aeromonasspecies are commonly isolated from fecal sample of children under the age of five years, whereas their isolation from other body sites usually occurred in adult populations. Aeromonads are known to cause severe diarrheal disease of short duration or chronic loose stools in children, the elderly, or the immuno-compromised individuals, and they have been implicated in travelers' diarrhea [4,5].

Tetracyclines belong to a family of broadspectrum antibiotics that includes tetracycline, chlortetracycline, doxycycline, and minocycline. These antibiotics inhibit protein synthesis in gram-positive and gram-negative bacteria by preventing the binding of aminoacyl-tRNA molecules to the 30S ribosomal subunit and inhibiting protein synthesis [6]. Contributing to higher levels of microbial resistance, especially among the genus *Aeromonas* [5].

This aim of this study was to evaluate the incidence and the spreading of A. *hydrophila* from diarrheic patients in Hilla city, Iraq, and study the antibiotic resistance patterns of the tested organisms to tetracycline and other antibiotics.

### <u>Material and Methods</u>

### **Collection of fecal samples**

This cross sectional study was designed to evaluate the incidence and the spreading of *A. hydrophila* from diarrheic patients in Hilla city A total of 822 fecal samples were collected. They were collected from rectal swab (routine work) and from patients suffering from diarrhea who attending public health lab, Hilla city, Iraq.

Specimen collection and analysis was carried out from October 2013 to February 2014.

# Isolation and identification of bacterial isolates

All specimens were cultured on alkaline peptone water, then transfer to TCBS and agarby MacConkey swabbing and incubated at 37°C for 24 hr. Each primary positive culture identified depending on the morphological properties such as (Shape, swarming, odor and lactose or non-lactose fermentation on MacConkey) [7]. Different Biochemical tests were used for identification of bacterial isolates according to standard methods [7, 8] The Vitek 2 svstem was used to confirm the biochemical according test to the manufacturer's instructions.

### **DNA Extraction and Purification**

A single colony of cultivated bacteria, which had been incubated overnight, transferred to 2 ml of sterile Louria broth and incubate at 37 °Cfor 18-20 hours. The DNA extracted and purified using Genomic DNA kit (EURx. /Poland Gene MATRIX). All clinical isolates were screened for chromosomal DNA according to manufacture instructions. The total DNA was used to detect16S rRNA. The DNA primers (*16SrRNA F*: CCAGCAGCCGCGGTAATACG,

*16SrRNA* R: TACCAGGGTATCTAATCC), 300 bp, were re-suspended by dissolving the lyophilized product after spinning down briefly with TE buffer molecular grad depending on manufacturer instruction as stock suspension. Working primer tube was prepared by diluted with TE buffer molecular grad.

# PCR thermocycling conditions and agarose gel electrophoresis:

The PCR tubes were placed on the PCR machine and the right PCR cycling program parameters conditions were as follows: 94° C 3min 1x, 94°C 30sec, 52° C 30sec 30x, 72° C, 30 sec, and 72° C 10 min. 1x.

The amplified PCR products were detected by agarose gel electrophoresis and visualized by staining with Ecodyes. The electrophoresis result was detected by using Biometra gel documentation system.

### Antimicrobial Susceptibility Testing

The antimicrobial susceptibility patterns of isolates to different antibiotics were determined using disk diffusion test (DDT) and interpreted according to CLSI guidelines [10]. The following antibiotics

were obtained (from Oxoid, UK, and Himedia, India) as standard reference disks as known potency for laboratory use: penicillin (P, 10 units), Piperacillin (PRL,100 amoxicillin-clavulanate μg), (AMC, 20/10 2g), Imipenem (IMP,10 μg), Meropenem (MEM, 10 µg), Gentamycin (CN,10 µg), Tobramycin (TM.5 µg), Amikacin Ceftazidime (AK10 μg), (CAZ,30µg), Cefotaxime (CTX,30 µg), Ciprofloxacin (CIP,10 µg), Tetracycline (TE,10µg), and Chloramphenicol (C, 30 μg)

**Determination of MICs of tetracycline** 

The ager dilution susceptibility method was used for determination of MICs of tetracycline according CLSI to documentations [10]. The ranges of appropriate dilutions of tetracycline MIC determination were 0.25-256 (µg/ml). To determine ager dilution break points, the plates were placed on dark surface, and the MIC was recorded as the lowest concentration of the antimicrobial agent that completely inhibits growth. MIC values were compared with the break points recommended by [10].

### **Results and Discussion**

### Isolation of Aeromonashydrophila

Results of this study revealed that out of 822 clinical sample 13 isolates (1.58%) were belonged to *Aeromonas*spp., however other bacterial isolates belong to other genera similar to *Aeromonas*were also recovered (Table 1).

Sample	Bacteria type	NO. of isolates	%
Positive	Aeromonas spp.	13	1.58%
Negative	Pseudomonas Spp., Pantoea spp. and Proteus Spp. and Enterobacter cloacae	809	98.4%
Total		822	100 %

<u>**Table 1:**</u>Occurrence of *Aeromonas* spp. recovered from fecal specimens

The low isolation rate of this bacteria 1.58% may be attributed to the fact that most cases of diseases caused by this bacteria are occurred usually in warm months, while the collection of samples in this study was in cold months. Aggeet al. reported that A. hydrophila similar to other enteric pathogens was seen more often in hot months. the frequency of A. hydrophila cases during warm months was 1.25 case per month, whereas during cold months it was 0.83 case per month [11]. The other reason of this result may be due to fact that A.hydrophila infects mainly children, elderly, and immunocompromised persons, while this study was focused on subjects of youth age group. Most studies are focusing isolating bacteria from feces of children only, and the fact that bacteria Aeromonas occur in children under two years at high rates, because of lack immune system completely, and abase infant formula of milk plays role in promoting the growth and reproduction of bacteria [12].

In a local study, Obaid [13] reported that 2.7% of A. hydrophila isolates were recovered from 479 patients from different ages and sexes. Naji [14] isolated this bacteria from children, the isolation rate of A. hydrophila was 4.08%. However, several authors found higher isolation rate of Aeromonas from clinical cases. AL-Fathlawy [15]obtained 20.17% ofA. hydrophila from clinical and environmental sample. On contrast, Borchardtet al [16] showed low isolation rate (0.66%) of A.hydrophila among 2565 diarrheic stool specimens submitted to a Wisconcin clinical reference laboratory.

Results showed that (8) isolates were diagnosed as *A. hydrophila* (61.53%), while the other isolates were distributed as (4, 30.7%) *A. salmoncidia* and (1, 7.6%) *A. sobria.* 

Result of isolation rate in the present study was similar to many studies conducted worldwide, Kannan *et al.* [17] found that the isolation rates of *A. hydrophila* were 60%, and 58.8% respectively, also they found that several species of *Aeromonas* were detected from acute diarrhea which were *A. caviae* (20%), *A. veronii* (10%), *A. schubertii* (4%), *A. jandaei* (3%), and *A. trota* (3%).

*A. hydrophila* and *A. sobria* tended to cause acute infection in human [18], while *A. salmonicida* cannot grow at 37°C, it is not pathogenic to humans [19]. Authors also referred to isolate *Aeromonas* spp. from different clinical specimens like blood (63%), wounds (11%), ascites (9%), feces (8%), and bile (3%). In addition to different unknown body sites [20].

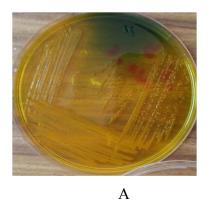
### Identification of A. hydrophila

Members of the genus *Aeromonas* are not difficult to isolate from clinical specimens in the diagnostic laboratory, but are often misidentified as belonging to the genus *Vibrio* or *Plesiomonas*[20]. Results of the phenotypic characteristics of the colonies *Aeromonas* had shown conformity with that reported by several authors [21, 22,23]. Bacterial isolation showed a good growth of *A. hydrophila* on TCBS medium and isolates produced yellow colonies /green color due to sucrose fermentation, with diameter of colonies ranged from 2-3 mm, while on blood agar, colonies appeared dark grey color beta- hemolytic (Figure 1).

On the MacConkey agar formed relatively small pale colonies is non-lactose fermenter (Table 2). also *A. hydrophila*  showed good growth in anaerobic condition because, *Aeromonashydrophila* facultative anaerobic, that is known to be pathogenic in humans [24].

Table 2: Characters of A.	hydrophilaisolates on	different culture media
	nyur opnituisoittes on	uniterent culture mould

Medium	Characters of colonies
TCBS agar	yellow shin with diameter ranged from (2-3)mm.
MacConkey agar	As pale like shaped indicated that <i>A.hydrophila</i> is unable to ferment lactose sugar
Blood agar	smooth, convex, rounded and $\beta$ -hemolytic colonies and pale white to grey color



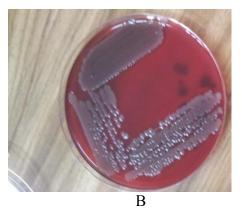


Figure 1: Characters of Aeromonashydrophila on (A) TCBS agar, (B) Blood agar

The microscopic examination of the bacteria stained by gram stain showed that the cells were gram negative, rod shaped, and the cells appeared singly to pairs, or as short chains [25].

Results ofbiochemical tests carried out for identification of isolates were compared with standard methods [10,1]. All isolates were positive for oxidase, and catalase .Oxidase test is used for differential of A. hvdrophila from other enteric bacteria. Results also found that A. hydrophila isolates had the ability to ferment glucose on Kligler iron ager (Alk/acid). They appeared positive to heamolysis test, motility test, and utilization of citrate, but they were negative to string test and urease test. String test is used to differentiate between A.hydrophila and V. cholera isolates [26].

Identification of *A. hydrophila* was confirmed using Vitek 2 system. Out of 13 *A.hydrophila*isolates (identified using biochemical tests), only 8 isolates was identified as *A.hydrophila*. The other 5 isolates were identified as *A. soberi* (1 isolate) and *A. salamoncidia* (4 isolates) had showed 85- 99% identification percentage probability.

**Molecular identification of** *A. hydrophila* 16S rRNA fragment was used for molecular identification of *A.hydrophila* isolates. Results found that 16S rRNA gene showed 100% similarity with *A. hydrophila* (Figure 3)that were identified previously as *A. hydrophila* using Vitek 2 system.

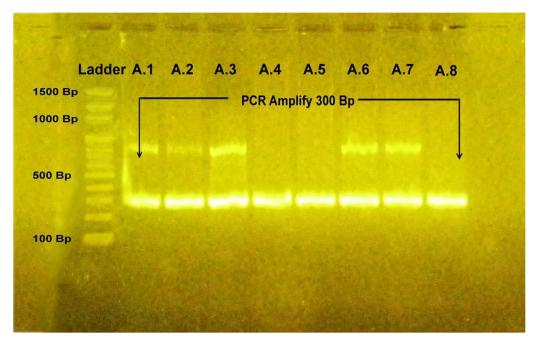
Identification of *Aeromonas* to the species level is difficult and complex due to their phenotypic and genotypic heterogeneity. The 16S rRNA ribosomal PCR amplifiedproduct size was 300 bp. that selected specific primer to this gene according to Jun *et al.*, [27]. The difficulty in identifying *Aeromonas* to the level of species was solved through diagnoses by viteks 2 system especially between *A. hydrophila A. cavia.* However, the molecular identification of isolates confirmed that isolate 8 of *Aeromonas* spp. isolates were belonged to *A.hydrophila.* 

The first attempts to identify *Aeromonas*genotypically relied upon differences in16S ribosomal DNA sequences was by Martinez-Murcia et al. [28], and several investigators developed for detection of probes various Aeromonasspp. [28, 29]. Several authors referred that 16rRNA gene was a specific

and a good marker in identification of all strains of *A.hydrophila* [20,30, 14].

No product was detected when genomic DNA from organisms other than *A. hydrophila* was used. 16S rRNA is a significant target to the molecular level identification. The upstream region of 16S rRNA is known to be highly conserved in species to species so this region could also be used for the verification of the thermodynamic stability on the basis of conserved secondary structures of RNA.

Different sources (other than 16S rRNA) of *A. hydrophila*have also been detected by the amplification of aerolysin gene [31], which targets 209 bp fragment of *aero* gene coding for the aerolysin toxin.



**Figure 2:**Ecodye stained agarose gel electrophoresis (1.2 %) of PCR amplified of *16 SRNA* gene (300) bp of *A. hydrophila* isolates

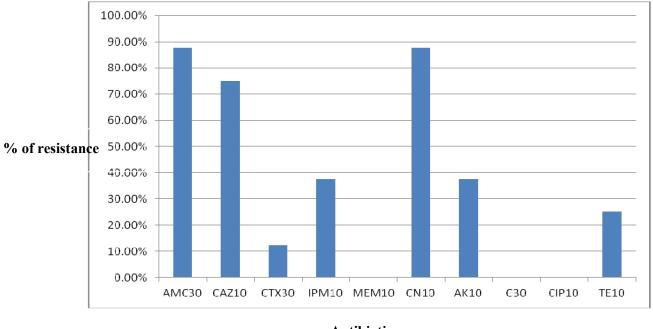
### Antimicrobial susceptibility testing

Results of susceptibility testing using DDT of *A. hydrophila* isolates (No.= 8) showed that 5 isolates (62.5%) were MDR. The definition of antibiotic resistance patterns was determined according to [36] who defined the MDR phenotype as resistance to representative antimicrobial agents of at least 3 different classes of drugs. Only three isolates (S2, S3, and S5,) were sensitive to most of classes of antibiotics (Table 3). However, no isolate showed XDR or PDR pattern of resistance. Results also showed that most isolates (87.5%) were resistant to amoxicillin-clavulanic acid and gentamycin. Six of *A. hydrophila*isolates (75%)were resistant to ceftazidime, whereas more than half of them were susceptible to amikacin, cefotaxime, and tetracycline. However 3 isolates (37.5%) were highly susceptible

(100%) to imipenem, meropenem and chloramphenicol (Figure 3).

Isolate	AMC	CAZ	СТХ	IPM	MEM	CN	AK	С	CIP	TE	Resistance Pattern
S1	R	R	R	R	S	R	S	S	S	S	MDR
S2	R	S	S	S	S	R	R	S	S	S	Sensitive
S3	R	S	S	S	S	R	R	S	S	S	Sensitive
S4	R	R	S	S	S	R	S	S	S	S	MDR
S5	S	R	S	S	S	S	S	S	S	S	Sensitive
S6	R	R	R	S	S	R	S	S	S	S	MDR
S7	R	R	R	S	S	R	R	S	S	R	MDR
S8	R	R	R	S	S	R	R	S	S	R	MDR
% of											
resistance	87.5	75	12.5	37.5	0	87.5	37.5	0	0	25	

Table 3: Antibiotic susce	ptibility patterns	of Aeromonashyd	<i>rophila</i> isolates



Antibiotic

Figure 3: Antibiotic susceptibility testing of A. hydrophila isolates using DDT

Absolute resistance of *Aeromonas*to ampicillin and oxacillin [33]. However, in this study susceptibility of *A. hydrophila* isolated were not tested for these antibiotic due to that they are not included in CLSI

documentations released from 2005 through 2014 [10, 34].

Several authors [37,38] found a similar findings regarding susceptibility to ceftazidime, cefotaxime, ciprofloxacin, and chloramphenicol. They revealed that *A*. *hydrophilia* isolated from Alice showed susceptibility to these antibiotics at 99%, 100%, 83.3%, and 83.3% respectively.

*Aeromonas* species are slightly susceptible to gentamycin. The aminoglycosides (amikacin, gentamicin, and tobramycin) showed excellent activity against almost all the isolates of the aeromonads except a few isolates of *A. cavie*[37].

Ashiru et al, [38] showed that all the species of Aeromonas (A. caviae, A. sobria and A. hydrophila) isolated from were resistant to nitrofurantoin and augmentin and randomly sensitive to ceftriazone, gentamycin, cotrimozazole and amoxicillin. Abulhamd [39] reported that a total of 10 motile Aeromonas strains were detected in water samples, Antimicrobial sensitivity patterns of the Aeromonas isolates revealed that 100% were sensitive to gentamicin, 80% to sulpha-methoxazole-trimethoprim, chloramphenicol. 70% to 50% to ciprofloxacin, 40% to neomycin, (30% to tetracycline, 20% to streptomycin and 10% to erythromycin.

Regarding to tet resistance, Ashiru*et al.* [40] revealed that *A. caviae*, *A. sobria*, and *A. hydrophila* isolated from water treatment were all resistant to tetracycline. The resistance to tetracycline has been reported to be acquired and encoded by plasmids or

transposons. Tetracycline inhibition have been reported to give excellent activity against the Aeromonads [41].

Other studies showed Antimicrobial susceptibility testing of the bacterial isolates using tetracycline discs demonstrated strong resistance to i.e., tetracycline in several isolates. Aeromonas Citrobacterfreundi, spp., Yersinia ruckeri, Pseudomonas putida [42].

Antibiotic resistance frequencies and profile varied according to the source of the strains. In this sense, one isolate exhibited resistance to seven antibiotics including three aminoglycosides, tetracycline, chloramphenicol, and trimethoprim/sulfa methoxazole), two were resistant to streptomycin, two were resistant to four aminoglycosides, and three were resistant to tetracycline and trimethoprim/ sulfa methoxazole[43].

### MIC of *A.hydrophila*isolates

Results of MIC of tetracycline for *A. hydrophila* isolates (NO= 8) found that only 2 isolates (25 %) were resistant to tetracycline (Table 4). The MIC of these isolates ranged from 0.125-16  $\mu$ g/ml. This result confirms tetracycline resistance by these isolates when tested using DDT (Table 3).

Isolates NO.	MIC of tetracycline (≥ 16 µg / ml)
S1	0.125
S2	0.125
S3	0.125
S4	0.125
S5	0.125
S6	0.125
S7	≥16
S8	≥16

Table 4: Determination of MIC of A.hydrophilaisolates

Changhesh *et al.* (2013) [44] recorded that the percentage of *A. hydrophila* isolated from diarrheic children (n=22) to tetracycline was 18.2%. Ko*et al.* [20] reported that fifty-one of all isolates of *A. hydrophila* from blood were susceptible to tetracycline.

Fass and Barnishan [46]carried out the MIC of 32 antimicrobial agents for 20 strains of A. hydrophila using by microdilution method and they found that the MIC values of tetracycline ranged from 0.5 -2  $\mu$ g/ml. They also showed among the other antimicrobial studied, agents only tetracycline. chloramphenicol, and trimethoprim-sulfamethoxazole were consistently active.

Change and Bolton [47] proved differences in resistance patterns were observed between strains isolated from different geographic locations and between *A. sobria* and *A. hydrophila* isolates. They also found that susceptibility to tetracycline was high (94.36%), consistent with previous reports from Australia and the United States.

Two local studies conducted in Iraq, found that all *A. hydrophila* isolated from clinical and environmental sources had 100% sensitivity to tetracycline when tested by DDT [14,15].

### **Conclusion**

The important conclusions in the present work can be summarized in the following points: Aeromonashydrophila was predominant among other species of isolation Aeromonasand the of Α. salmoncidia from human specimens in the present study represented as a first record in Iraq. the results of susceptibility testing using DDT of A. hydrophila isolates showed that five isolates were MDR, Only three isolates were sensitive to most of classes of antibiotics. Results of MIC found that only 2 isolates were resistant to tetracycline.

### **References**

1. ErdemB., Kariptas E., Cil E., and Isik K. (2011). Biochemical identification and numerical taxonomy of *Aeromonas* spp. isolated from food samples in Turkey. A thesis of Science, Artsand Science Faculty, Evraw Univ., 35: 463 -472.

2. Sarkar A., Saha M. and Roy P. R. (2012). Identification and typing of *Aeromonashydrophila* through 16 s DNA – PCR finger printing. J. Aquacult. Res. Der 3:6.

3. Abbott S.L., Cheung W.K.W., and Janda J.M. (2011). The Genus *Aeromonas*: Biochemical characteristics, Atpyical Reactions, and phenotypic Identification Schemes. J. Clin. Microbiol., 41 (6): 2348-2357.

4. Igbinosy I.H., Tom M., Ignmbor E.U., Okoh A.; and Ighdasi F. (2012). Emerging *Aeromonas* species infections and their significance in public Health. The Scientific world J. doi:10.1100 /2012/ 625023.

5. Markov G., Kirov G., Lyutskanoy V., and Kondarer M. (2007). NocrotizingFascitis and Myonecrosis due to *Aeromonashydrophila*. Wounds,19(8): 223-226.

6. Aminov, R.I., Garrigues-Jeanjean, N. and Mackie, R.I. (2001). Molecular ecology of tetracycline resistance: development and validation of primers for detection of tetracycline resistance genes encoding ribosomal protection proteins. Applied and Environmental Microbiology 67, 22–32.

7. Brenner D.J., Krieg, N.R. and Staley, J.S. (2005). Bergeys manual of systematic bacteriology. vol 2,2<sup>nd</sup> edition. partB, pp.556-578.

8. Forbes, B.A., Daniel, F.S. and Alice, S. W. (2007). Bailey and Scotts diagnostic microbiology .12 th ed., Mosby Elsevier company, USA.

9. MacFaddin, J.K. (2000). Biochemical test for identification of medical bacteria. 3<sup>rd</sup> ed. Lippincott Williams and Winkins. A Wolter Kluwer Company. Philadelphia Baltimor. New York.

10. Clinical and Laboratory standards institute (CLSI). (2005). Performance standards for antimicrobial susceptibility testing. Twenty third informational supplement.

11. Aggeri W.A., McCormick J.D., and Gurwith M.J. (1985). Clinical and Microbiological feature of *Aeromonashydrophila* associated diarrhea. J. Clin. Microbiol., 909-913.

12. Wong H. (2007) Enteropathogenic Aeromonads. Department of Microbiology, Soochow University.

13. Obaid, A.J. (2013). S-Layer role of *Aeromonashydrophila* isolated from diarrhea specimens. M.Sc thesis, College of Science, University of Babylon.

14. Naji, N.Q. (2013). Dissemination and Immunological properities of *Aeromonashydrophila*isolates from children with diarrhea in AL-Muthana province. M.Sc thesis, College of Science, University of AL-Muthanna.

15. AL-Fatlawy H.N.K. (2012). Bacteriological and Molecular study of *Aeromonashydrophila* isolated from clinical and Environmental samples. M.Sc thesis, College of Science, University of Kufa.

16. AL-Fatlawy H.N.K. (2012). Bacteriological and Molecular study of *Aeromonashydrophila* isolated from clinical and Environmental samples. M.Sc thesis, College of Science, University of Kufa.

17. Borchardt M. A., Stenper M. E., and Standridge J.H. (2003). *Aeromonas* isolates from Human Diarrheic stool and Groundwater compared by pulsed–field gel electrophoresis. Emerging infectious Diseases, 9 (9).

18. Moyer N.P.(1987).Clinical significance of *Aeromonas* species isolated from patients with diarrhea. J. Clin. Microbiol., 2044-2048.

19. Altwegg, M., A. G. Steigerwalt, R. Altwegg-Bissig, J. Lüthy-Hottenstein, and

D. Brenner. (1990). Biochemical identification of *Aeromonas*geno-species isolated from humans. J. Clin. Microbiol., 28: 258-264.

20. Ko w .C, Yu K.W, Liu C.Y(1996). Increasing antibiotic resistancein clinical isolates of *Aeromonas* strains in Taiwan. Antimicrob Agents Chemother,40:1260

21. Puthucheary S.D.; Puah S.M.; and Chua K.H. (2012). Molecular characterization of clinical isolates of *Aeromonas* species from Malaysia. Plos one,7(2): e30205. Doi: 10.1371.

22. JandaJ.M.A., and Abbott S.L. (2011). The genus *Aeromonas*:Taxonomy, pathogenicity, and infection. Clin Microbial Rev., 23(1): 35-73.

23. Chu, W.H and Lu, C.P. (2005). Multiplex PCR assayfor the detection of pathogenic *Aeromonashydrophila*. Journal of Fish Diseases, 28: 437–441.

24. Okumura R., Liao C., Gavin M., Jacoby G. A., and Hooper D.C. (2011). Quinolone induction of qnr VSI in *Vibrio splendidus* and plasmid – carried qnrSI in *Escherichia coli*, mechanism independent of the SOS system. Antimicrob. agents Chemother., 55(12): 5942-5945.

25. Villari, P., Crispino, M., Montuori, P., and Boccia, S. (2003) Molecular typing of *Aeromonas* isolates in natural mineral waters. App. Environmen. Microbiol., 69: 697-701.

26. Obi, C. L., Ramalivhana, J., Samie, A., and Igumbo, E.O. (2007). Prevalence, pathogenesis, antibiotic susceptibility profiles, and *in vitro* activity of selected medicinal plants against *Aeromonas* isolates from stool samples of patients in the Venda region of South Africa. J. Health Popul. Nutr. 25(4):428-35.

27. Perilla, M.J.; Ajello,G.; Boop,C.; Elliot,J.; Facklan,R.; Popovic,T. and Wells, J. (2003). Manual for laboratory identification and antimicrobial susceptibility testing of bacterial pathogens of public health importance in developing world, Center for Diseases Control and Prevention, Atlanta, Georgia, USA.

Jun. J.W.; Kim, J.H.; Gomez, D.K.; Choresca Jr, C.H.; Han, J.E.; Shin, S.P., and Park. S.C. (2010). Occurrence of 36. resistant

tetracycline Aeromonashvdrophilainfection in Korean

28.

cyprinid loach (Misgurnusanguillicaudatus). Afr. J. Microbiol. Res., 4(9): 849-855.

Martinez-Murcia, A. J., Benlloch, S. 29. and Collins, M. D. (1992). Phylogenetic interrelation-ships of members of the Aeromonasand Plesiomonasas genera determined by 16S ribosomal DNA sequencing: lack of congruence with results of DNA-DNA hybridizations. Int. J. Syst. Bacteriol. 42:412–421.

30. Demarta, A., M. Tonolla, A. Caminada, M. Beretta, and R. Peduzzi (2008).Epidemiological relationships between Aeromonasstrains isolated from children and symptomatic household environments as determined by ribotyping. European Journal of Epidemiology 16(5): 447-453.

Khan A.A. Cerniglia C.E. (1999). 31. Rapid and sensitive method for the detection *Aeromonascaviae* and of *Aeromonastrotaby* polymerase chain reaction. Letters App. Microbiol., 24:233-239.

32. Chang C.I., Hung, P.H., Wu, C., Cheng, T.C.; Lin, K.J. (2012). Simultaneous Detection of Multiple Fish Pathogens Using a Naked-Eye Readable DNA Microarray.Sensors, (12): 2710-2728. 33. Al-Fatlawy H.N.K.; and AL-Ammar M.H. (2013). Molecular study of Aeromonashydrophila isolated from stool samples in Najaf (Iraq). Int. J. Microb. Res., 1:363-366.

34. Clinical and Laboratory standards institute (CLSI). (2014). Performance standards for antimicrobial susceptibility testing. Twenty fourth informational supplement.

35. Pollard, D.R.; Johnson, M.W.; Lior, H.; Tyler, S.D. and; Rozee, K.R. (1990). Detection of the aerolysin gene in

Aeromonashydrophila by PCR. J Clin. Microbial.82:2477-2481.

Magiorakos A., Srinivasan P., Carey R.B., Carmeli Y., Falagas M.E., Giske C.G., Harbath S., HindlerJ.F., Kahlmeter G.,Olsson-Lilijequist B., Paterson D.L., Rice L.B., Stelling, M.J., Struelens M.J., Vatopoulos A., Weber J.T., and Monnet D.L. (2012). Multidrugresistant, extensively drug-resistant and pan-drug resistant bacteria: an international expert proposal for interim standard definitions for acquired resistance. Clin Microbial. Infect 18:268-281.

Igbinosy I.H., Tom M., Ignmbor 37. E.U., Okoh A., and Ighdasi F. (2012). Emerging Aeromonas species infections and their significance in public Health. The Scientific world J. doi :10.1100 /2012/ 625023.

38. Dias C., Mota V., Martinez-Murcia A., and Saavedra M.J. (2012).Antimicrobial resistance patterns of Aeromonas spp. isolated from Oranamental fish. J. Aquacult Res. Der 3:3.

Awan M. B., Magbool A., Bari A., 39. and Krovacek K. (2009). Antibiotic susceptibility profile of Aeromonas spp. isolates from food in Abu- Dhabi .United Arab Emirates. New Microbiol., 32: 17-23.

Ashira A.W., Uaboi – Egbeni P.O., 40. Oguntowo J.E., and Idika C. N. (2011). Isolation antibiotic and profile of Aeromonas species from Tilapia fish (*Tilapianilotica*) and cat fish (Clariasbetrachus). Pak. J. Nut., 10 (10): 982 - 986.

41. Abulhamd A. (2010). Genetic diversity and antimicrobial susceptibility of motile aquatic aeromonads. Int. J. Chemother. Engineer. Appl.,1(1): 2010 – 2021.

42. Vivekanandhan G., Savithamani K., Hatha.A.M., Lakshmanaperumalsamy P. (2002).Antibiotic resistance of Aeromonashydrophila isolated from marketed fish and prawn of South India. Int. J. Food Microbiol., 76 (2002) 165-168.

43. HedyatianfardM. and Sharifiyazdi H. (2014). Detection of tetracycline resistance genes in bacteria isolated from fish farms using polymerase chain reaction. veterinary Research forum 5(4): 269-272.

44. Guerra, I.M. F., Fadanelli-Raquel F.M., Schreiner F. D.; Anapaula L.; Wollheim C.C.; Sergio O. P. (2007). *Aeromonas* associated Diarrheal disease in south Brazil prevalence virulence factors and antimicrobial resistance. Braz. J. Microbiol., 38: 638-643.

45. Chenghesh K.S., EL-Mohammady H., Levin S.Y., Zorgani A., and Tawil K.

(2013). Antimicrobial resistance profile of *Aeromonas* species isolated from Libya. Libyan J. Med, 8:1-2.

46. Fass, R.J. and J. Barnishan (1981). *In vitro* susceptibility of *Aeromonashydrophila* to 32 antibiotics. Antimicrob. Agents Chemother., 19: 357-358.

47. Chang B.J., and Bolton S.M. (1987). Plasmids and resistance to antimicrobial agents in *Aeromonassobria* and *Aeromonashydrophila* clinical isolates. Antimicrob. Agents Chemother., 31(8): 1281-1282.