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Heavy Metal Resistance of Aeromonas hydrophila Isolated from Raw and **Drinking Water in Baghdad City**

Sanaa R. Oleiwi*

Department of Biology, College of Sciences, University of Baghdad, Baghdad, Iraq

Abstract

A total of 551 water samples (drinking and raw water) were collected In this study, Aeromonas.hydrophila, were detected by biochemical tests and PCR (16s rRNA gene). The results of identification showed that A.hydrophila had recovery rate 63 isolates (49.21%). The results revealed that all A.hydrophila isolates were PCR positive or the 16S rRNA gene and the results of sequencing showed that two isolates of A.hydrophila(local isolates) had percentage similarities 100% with A. hydrophila ATCC 7966 in GenBank database .All strains had a minimal Inhibitory Concentration(MIC) distribution pattern for lead cetate rranged (900-1200 µg/ml), and mercury chloride ranged (40-80 µg/ml).

Keywords: Heavy Metal, *Aeromonas hydrophila*, Drinking Water. Resistance.

مقاومة المعادن الثقيلة لبكتريا Aeromonas hydrophila المعزولة من المياة الخام ومياه الشرب في مدينة بغداد

سناء رحمن عليوي *

قسم علوم الحياة، كلية العلوم، جامعة بغداد، بغداد، العراق

الخلاصة

جمعت 551 عينة ماء خام وماء شرب وتم عزل بكتريا Aeromonas hydrophila وتم تشخيصها بوساطة الفحوصات الكيموحيوية وتقنية تفاعل البلمرة وتتابع النيوكليوتيدات .اظهرت نتائج التشخيص ان نسبة العزل كانت 49.21% (63 عزلة بكتيرية)وكذلك اعطت جميع عزلات البكتريا نتيجة موجبة لتفاعل البلمرة المتسلسل (تتابع النيوكليوتيدات) وكانت عزلتان محليتان ذات نسبة تشابة 100% مع العزلات القياسية في بنك الجينات اعطت جميع العزلات تراكيز المثبطة الدنيا التي تراوحت بين 900-1200 مايكروغرام/ مل لخلات الرصاص و 40-80 مايكروغرام/ مل لكلوريد الزئبق.

Introduction

Aeromonas hydrophila is a Gram negative bacteria widely distributed in freshwater environments [1]. It is a well-known fish [2, 3] and human pathogen [4]. Since some strains of Aeromonas are enteropathogens possessing a range of virulence factors (enterotoxins, cytotoxins, haemolysins, and invasive ability), infected fish may be vehicles of human infection [5-7] some are heavy metal resistant [8].

The presence of toxic heavy metals in the environment has resulted in the development or acquisition by bacteria of genetic systems that counteract their effects. Many bacterial heavy metal resistance systems are based on efflux, and two groups of efflux systems have been identified. These can be either P-type ATPases, e.g., the Cu(II), Cd(II), and Zn(II) ATPases of gram-negative bacteria [9].

^{*}Email: sanaeleiwi@yahoo.com

Material and methods

One hundred and thirty one raw water samples (from Tigris river) and four hundred and twenty drinking water samples from five water treatment plants (WTPs) in different area in Baghdad) were collected. Prior to collection of drinking water was allowed to run at a uniform rate for 2–3 min. in a sterile bottles containing sodium thiosulphate to a final concentration of 0.01% (W/V) to neutralize any free or combined residual chlorine. One liter of water sample was collected in each bottle, the samples were carried out to the laboratory special aspectic cool box. All the samples were transported to the laboratory on ice and analyzed within 24 h [10].

Drinking water were filtered by filtration apparatus then the membranes were then transfered carefully to a Petri dish of Ampicillin Dextrin Agar(ADA), then the Petri dishes were incubated at 37 °C for 24 hours. Aloop full of river water samples was cultured directly on Ampicillin Dextrine Agar(ADA) and incubated at 37 °C for 24 h[11].

Identification of bacterial isolates

Bacterial isolates were identified by the procedures described in Bergey's Manual of Systematic Bacteriology [12].

For further identification of the *Aeromonas spp.* isolates, two API strips: API 20 E, ID32 E, also these isolats were identified by PCR *16s r*RNA gene,by using Primers: 5-GGGAGTGCCTTCGGGAATCAGA-3(Forward primer) and 5-TCACCGCAACATTCTGATTTG-3(Reverse primer) [13].

Determination of Minimal Inhibitory Concentration (MIC) of heavy metals resistance

The minimal inhibitory concentration (MIC) of metals (Mercury chloride and Lead acetate) was determined by an agar dilution procedure, as previously described [8, 14-16]. A.hydrophila isolates grown on heavy metals incorporated media (nutrient agar pH $\,7$), Heavy metals concentrations used in this test were :900,1000,1100, and 1200 µg/ml for lead acetate and 40,50,60,70, and 80 µg/ml for mercury chloride.

MIC was observed when the isolates failed to grow on plates after incubation period at 37 °C for 24 hr , MIC breakpoints (μ g/ml) for considering a bacterial isolate as susceptible (S) or resistant (R) to lead and mercury as reported by [14] : lead S <800, R >800, mercury S <54, R >54.(S:sensitive ,R:resistance).

Results and Discussion

The results of identification showed that A. hydrophila had recovery rate 63 isolates (49.21%), forty five A. hydrophila isolates from raw water (Tigris river) and eighteen isolates from drinking water were obtained.

The results revealed that all *A. hydrophila* isolates were PCR positive for the 16S rRNA gene Figure-1,-2, and the results of sequencing showed that two isolates of *A. hydrophila*(local isolates) had percentage similarities 100% with *A. hydrophila* ATCC 7966 in GenBank database(according to sequencing were carried out in Korea using Sequencer applied biosystem (3730XL)USA (genetic analyzer 3730XL). Figure-3.

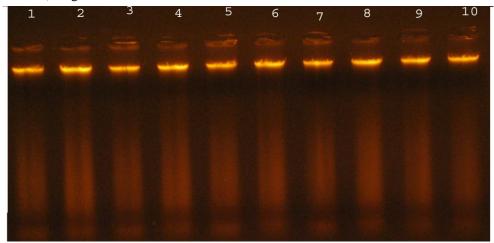


Figure 1-Agaros gel electrophoresis (1%) of genomic DNA of *A. hydrophila* isolates ,1.5 hour at 5 V/cm, stained with ethidum bromide and visualized on a UV transilluminator. Lane 1-10: Genomic DNA extracted from different *A. hydrophila* . isolates. Purity between (1.8-1.9)

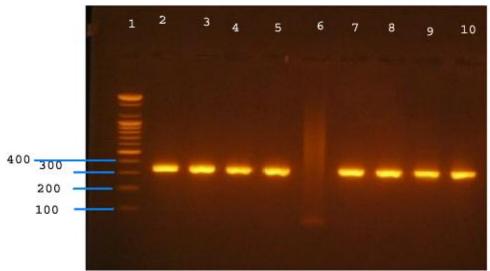


Figure 2-Agaros gel electrophoresis (1.5%) of amplified 16S rRNA gene (356 bp) of *A. hydrophila* for 1.5 hour at 5 V/cm, stained with ethidum bromide and visualized on a UV transilluminator.Lanes 1: 100 bp DNA ladder (2Kb). Lane 6: negative control. (had all PCR mixture except of DNAtemplate). Lane 2-5 and 7-10: *A. hydrophila* .

Alignment statistics for match #1

Score			Expect	Identities	Gaps	Strand
460 bits(249		9)	2e-126	249/249(100%)	0/249(0%)	Plus/Minus
Query	1	CTAGCTTGCA	GCCCTCTGT ACGCGC	CATT GT AGC AC GT GTG TAG CC CT GGC CGT	AAGGGC 60	
		пппппп	шшшшш		ШШ	
Sbjet	216174	CTAGCTTGCA	GCCCTCTGTACGCGC	CATT GT AGC ACGT GTG TAG CCCT GGC CGT	AAGGGC 216115	
Query	61	CATGATGACT	T GAC GT CAT CC CCA C	CTTCCTCCGGTTTATCACCGGCAGTCTCC	CTTGAG 120	
		1111111111			ШШ	
Sbjet	216114	CATGATGACT	T GAC GT CAT CC CCA C	CTTCCTCCGGTTTATCACCGGCAGTCTCC	CTTGAG 216055	
Query	121	TTCCCACCAT	TACGTGCTGGCAACA	AA GG AC AGG GG TT GCG CTC GT TG CGG GAC	ITAACC 180	
		ШШШ	111111111111111111111111111111111111111		111111	
Sbjet	216054	TTCCCACCAT	T ACGTG CTG GC AAC A	AA GG AC AGG GG TT GCG CTC GT TG CGG GAC	TTAACC 215995	
Query	181			ACAGCCATGCAGCACCTGTGTTCTGATTC		
Sbjet	215994	CAACATCTCA	CGACACGAGCTGACG	ACAG CCATG CA GC ACCTGT GT TC TGA TTC	CCGAAG 215935	
Query	241	GCACTCCCA	249			
		111111111				
Sbjet	215934	GCACTCCCA	215926			

Figure 3- PCR analysis of A. hydrophila isolates

Heavy metal resistance of A. hydrophila

All isolates had a (MIC) distribution pattern for lead acetate ranged (900-1200 $\mu g/ml$), and mercury chloride ranged (40-80 $\mu g/ml$). The results of MIC of heavy-metals (lead and mercury) are shown in the Table -1.

Table 1- MIC distribution patterns of heavy metals for *A.hydrophila*

Heavy metal	MIC (μg/ml)	% no. of inhibited isolates
	900	30.1%(19 isolates)
Load agetate (PhCH2COO)	1000	60.3% (38 isolates)
Lead acetate (PbCH3COO)	1100	87.3% (55 isolates)
	1200	100%
	40	9.5% (6 isolates)
	50	19.1%(12 isolates)
Mercury chlorid (HgCl2)	60	60.3%(38 isolates)
	70	80.9%(51 isolates)
	80	100%

MIC breakpoints ($\mu g/ml$) for considering a bacterial isolate as susceptible (S) or resistant (R) were those reported in [14] for lead S < 800, R >800; for mercury S <54, R >54.

All isolates (100%) of *A. hydrophila* were resistant to lead acetate that came in accordance with the finding of [15] and forty five isolates (71.42%) were resistant to mercury chloride that disagreement with results of Paniagua [15], who found that all isolates of *A.hydrophila* were susceptible to mercury chloride, that may due to different between isolates in current study and Paniagua. In study conducted by Miranda and Castillo (1998) [8] found that resistant aeromonads to heavy metals are easily recovered from water sources, posing a potential public health risk, that due to avariety of detoxifying mechanism developed by resistant microorganisms such as binding with bacterial cell envelopes, metal reduction, metal efflux [15].

Bacterial heavy metal resistance determinants commonly reside on plasmids, and selective pressure for one resistance indirectly leads to selection for the others, as the responsible genes are linked. Since many low-statuses farmer people consume these fish and usually swim in this freshwater pond, contact with the heavy metal resistant *Aeromonas* posess an important human health risk. In addition, passage of these bacteria from fish to humans would permit the horizontal gene transfer from *Aeromonas* to other bacteria harbored by the exposed persons[15, 16].

Reference

- **1.** Holmes P., Nicolls L.M. and Sartory D. P. **1996**. The ecology of mesophilic *Aeromonas* in the Aquatic environment. In: Austin, B., M. Altwegg, P.J. Gosling & S. Joseph (Eds.) *The genus Aeromonas*, John Wiley & Sons, New York, pp:127–150.
- **2.** Joseph S.W. and Carnahan, A. **1994** .The isolation, identification, and systematics of the motile *Aeromonas* species. *Annual Review of Fish Diseases*, 4, pp. 315–343.
- **3.** Austin Band Adams C.**1996**. Fish pathogens. *In*: Austin, B., M. Altwegg, P.J. Gosling, & S. Joseph (Eds.). *The genus Aeromonas*, John Wiley & Sons, New York, pp:197–243.
- **4.** Altwegg M. and Geiss, H.K. **1989**. *Aeromonas* as a human pathogen. *Critical Reviews in Microbiol*.,16, pp: 253–286.
- **5.** Cahill, M.M .**1990**. Virulence factors in motile *Aeromonas* species. *J.of Appl . Bacteriol.*, 69, pp:1-16.
- **6.** Kirov, S.M. **1993**. The public health significance of *Aeromonas spp*. In foods. *Inter. J. of Food Microbiol.*, 20, pp: 179-198.
- **7.** Muslim, S.N. **2005**. Biochemical and Genetic Studies on Hemolysin Produced by *Aeromonas hydrophila* Isolated from Diarrheal Patients and from Surface Water. A Ph.D. Thesis. College of Science, Al-Mustansiriyah University.
- **8.** Miranda, C. D. and Castillo, G. **1998**. Resistance to antibiotic and heavy metals of motile aeromonads from Chilean freshwater. *The Science of the Total Environment*, 224, pp. 167-176.
- 9. Silver S 1996. Bacterial resistances to toxic metal ions--a review. Gene. 7, 179(1), pp:9-19.

- **10.** Eaton A., Wef E. and Arnold E. **2005**. *Standered method for the examination of water and waste water*. Twenty First Edition. American public health association.
- **11.** Muhammad, B.A. **2005** .A comparative study of *Aeromonas hydrophila* isolated from local waters and clinical samples. M.Sc. Thesis, College of Science, University of Baghdad.
- **12.** Martin-Carnahan, A and Joseph S W. **2005**. *Aeromonadaceae*. In Brenner, D. J., N. R. Krieg, J. T. Staley, and G. M. Garrity. *The Proteobacteria, Part B, Bergey's Manual of Systematic Bacteriology*, Second Edition, Volume 2, Springer-Verlag, New York, NY.
- **13.** Wang, G., Clifford, C.G., liu, C., Pucknell, C., Munro, C.K., Kruk, T.M., Caldeira, R., Woodward, D.L. and Rodgers, F.G. **2003**. Detection and characterization of a hemolysin gene in *Aeromonas hydrophila* and *Aeromonas sobria* by multiplex PCR. *J. Clin. Microbiol.* ,41, pp: 1048-1054.
- **14.** Vaca, S., Mirandi, R. and Cervantesc. **1995**. Inorganic-ion resistance by bacteria isolated from a Mexico City freeway. *Antonie van Leeuwenhoek*, 67, pp. 333-337.
- **15.** Paniaguagl, Monroy E., Perches M., Negrete E. and vaca oGS. **2006** Antibiotic and heavy metal resistance of *Aeromonas hydrophila* isolated from charal (*Chirostoma humboldtianum*). *Hidrobiológica*, 16 (1), pp: 75-80.
- **16.** Shakooria.R, Zaidi,k.S.and Haq, R .**1998.**Cadmium resistance Enterobacterclacae and *Klebsiella* sp isolated from industrial effluents and their possible role in cadmium detoxification .*World J. of Microbiol.and biotechnol.*,15, pp:249-254.