

Aeromonas and *Pseudomonas*: antibiotic and heavy metal resistance species from Iskenderun Bay, Turkey (northeast Mediterranean Sea)

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Abstract We studied the susceptibility patterns to 15 different antibiotics and six heavy metals in *Aeromonas* spp. and *Pseudomonas* spp. isolated from Iskenderun Bay, Turkey (northeast Mediterranean Sea). A high percentage of *Aeromonas* isolates showed resistance to cefazolin (66.6%) and trimethoprim–sulphamethoxazole (66.6%). Amongst the *Pseudomonas* isolates, there was a high incidence of resistance to nitrofurantoin (86.2%), cefazolin (84.8%) and cefuroxime (71.7%). Most isolates showed tolerance to different concentrations of heavy metals, and minimal inhibition concentrations ranged from 25 to >3,200 µg/ml. The *Aeromonas* spp. and *Pseudomonas* spp. showed high resistance to copper of 98.3% and 75.4%, respectively, and low resistance to lead of 1.7% and 7.2%, respectively. Our results show that antibiotic and heavy metal resistant *Aeromonas* spp. and *Pseudomonas* spp. were widespread in Iskenderun Bay in 2007 and 2008. The increasing presence of antibiotic

and heavy metal resistant *Aeromonas* spp. and *Pseudomonas* spp. may become a potential human health hazard.

Keywords *Aeromonas* · *Pseudomonas* · Antibiotic resistance · Heavy metal resistance · Human health

Introduction

Aquatic bacteria that are resistant to multiple antibiotics are of great importance in many areas of the world. Many aquatic bacteria are responsible for various types of serious diseases; for this reason, they have been the focus of numerous studies (Calomiris et al. 1984; Messi et al. 2005; Lobo et al. 2008; Matyar et al. 2008). Hospitals discharge large quantities of untreated antibiotic wastes into the environment, which has led to an increase in bacteria having multiple antibiotic resistances (MARs) and to an increase in more virulent pathogens.

Aeromonas spp., autochthonous inhabitants of aquatic environments, are widespread in natural habitats such as soil, fresh and brackish water, sewage and wastewater (Araoju et al. 1991). These bacteria cause both gastrointestinal and, to a lesser extent, extra-intestinal infections in humans (Janda and Abbott 1998).

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Amongst the 14 species of *Aeromonas* known to date (Abbott et al. 2003), *Aeromonas hydrophila*, *Aeromonas caviae* and *Aeromonas veronii* biotype *sobria* have most commonly known in human infections and produce a variety of virulence factors such as haemolysins, cytotoxins, enterotoxins, proteases, leukocidin, phospholipases, endotoxins, outer membrane proteins and fimbriae (Chopra and Houston 1999). Pseudomonads are a ubiquitous group of environmental Gram-negative bacteria, amongst which *Pseudomonas aeruginosa* is the third most common pathogen responsible for hospital-acquired (nosocomial) infections (Schaberg et al. 1991).

Fluvial waters receive human and animal wastewater discharges, which are expected to contain anti-microbial agents likely to exert a selective pressure and commensally resistant bacteria capable of transferring their resistances to autochthonous bacteria (Goñi-Urriza et al. 2000). Although the antibiotic sensitivity of food and clinical isolates of *Aeromonas* spp. have been extensively studied (Monteil et al. 2004; Demerta et al. 2004; Palú et al. 2006; Wu et al. 2007), there is limited information about the presence of aeromonads in the marine environment.

Industrial and agricultural activities have led to substantial release of toxic metals in the environment, which can constitute a major hazard for ecosystem as well as human health (Nriagu and Pacyna 1988). In many aquatic systems, trace metals are significant contaminants, due in part to anthropogenic sources such as industry and mines inputs. Coal-fired power plants generate large quantities of ash through combustion, and this waste contains residues of multiple trace metals such as arsenic and selenium. The ash residue mixed with the water before being deposited in settling basins and then discharge into aquatic system (Rowe et al. 2002).

Ecological studies have reported that antibiotic and metal resistances are becoming a global importance (Hassen et al. 1998; Benka-Coker and Ekundayo 1998; Ansari and Malik 2007). Plasmids are known to carry resistance to antibiotics and heavy metals (Smith et al. 1993; Sobecky 1999). Marine bacteria adsorb, accumulate and transform heavy metals in the most food chains (Chan and Dean 1988).

We focused our present studies on Iskenderun Bay on the south coast of Turkey, lying in northern east of Mediterranean Sea (36°38' E, 36°05' N). Domestic wastes, including hospital wastes, are discharge into the bay. In addition, factories, like iron and steel, a fertiliser factory, a refinery and a coal-fired power plant discharge a high amount of processed or unprocessed wastes into the bay. Despite these environmental pressures, the bay is an important region for fishing and aquaculture. Recent studies have shown that there are high levels of accumulated heavy metals in fish from this bay (Türkmen et al. 2006; Atli and Canli 2007).

To the best of our knowledge, this is the first research determining the presence as well as resistance to antibiotics and heavy metals of *Aeromonas* spp. and *Pseudomonas* spp. isolated from Iskenderun Bay. The specific aims of this study were (a) to identify the *Aeromonas* spp. and *Pseudomonas* spp. strains recovered in Iskenderun Bay, (b) to determine the level of antibiotic resistance rates against widely used antibiotics in Turkey, (c) to determine the heavy metal resistance of the bacteria and (d) to investigate relationship between the antibiotic and heavy metal resistance (if present).

Materials and methods

Sampling

Our studies were conducted in three different sites along the southeast coast of Iskenderun Bay, Turkey which is located in the northeast of the Mediterranean Sea (Fig. 1). The geographic coordinates of the sampling sites were 1:35°56' E, 36°28' N; 2:36°11' E, 36°36' N and 3:36°10' E, 36°42' N.

Water samples (250 ml) were collected from March 2007 to February 2008. Samples were collected into sterile bacteriological sample bottles and brought to the laboratory in an ice chest (APHA 1992). The samples were processed within 4 h of collection. A total of 40 samples (13, 12 and 15 from sites 1, 2 and 3, respectively) were examined for the presence of *Aeromonas* spp. and *Pseudomonas* spp.

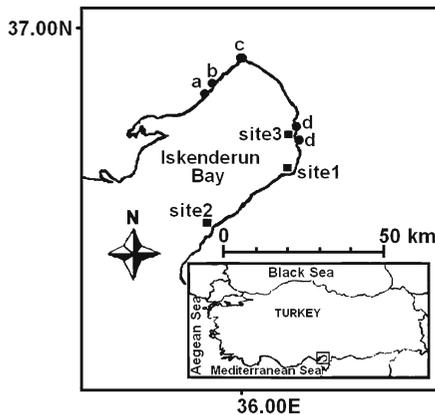


Fig. 1 Map of Turkey and Iskenderun Bay showing the three different sampling sites; *a* coal-fired power plant, *b* oil refinery, *c* fertiliser factory, *d* iron-steel factory

Bacterial isolation and identification

For isolation of *Aeromonas* and *Pseudomonas* spp., 25 ml of each water samples were inoculated in 225 ml alkaline pepton water (pH 8.6) with 1% of NaCl (*w/v*) and incubated at 30°C for 24 h. The samples were then plated onto selective media (ampicillin dextrin agar (ADA) medium and *Pseudomonas* F agar (Becton and Dickinson, Sparks Glencoe, MD, USA)) and then incubated at 30°C for 24–72 h. The ADA medium was freshly prepared having the following composition (grams per liter) according to Havelaar et al. (1987): tryptose (Difco), 5; dextrin (Merck), 10; yeast extract (Merck), 2; sodium chloride (Merck), 3; potassium chloride (Merck), 2; magnesium sulphate.7H₂O (Merck), 0.2; ferric chloride.6H₂O (Merck), 0.1; sodium desoxycholate (Merck), 0.1; bromothymol blue (Merck), 0.04; agar (Oxoid), 15 and ampicillin (ICN Biomedicals), 0.01, with pH of 8.0.

Presumptive *Aeromonas* spp. isolates were obtained from ADA medium by colony morphology (yellow, circular, convex, 1–3 mm diameter colonies). The genus *Aeromonas* was identified based on the findings of Gram-negative rods, positive oxidase test, fermentation of D-glucose, motile and the absence of growth in 6.5% sodium chloride (Ko et al. 2000).

Isolates were considered to be presumptive *Pseudomonas* spp. if they were Gram-negative

rods that produced fluorescent colonies on *Pseudomonas* F agar, were oxidase- and catalase-positive and were glucose oxidative and positive for citrate. Presumptive *Aeromonas* and *Pseudomonas* colonies were subcultured on tryptone soy agar (Oxoid) at 30°C for 24 h to obtain at least two consecutive pure cultures. Identity was confirmed by using the Becton and Dickinson Crystal E/NF ID system (BBL, Cockeysville, MD, USA). These strains were identified by using the E/NF identification software (BBL, Cockeysville, MD, USA). A number of isolates (20%) were reexamined to check the test reproducibility.

Test for antibiotic resistance

Antibiotic resistance was determined by an agar diffusion test (NCCLS 1997) using Mueller–Hinton agar (Difco). Fifteen different antibiotics (representing eight classes) were used. The antibiotics tested and their sensidisk concentrations were amikacin (AN; 30 µg), streptomycin (S; 10 µg), gentamicin (GM; 10 µg), kanamycin (K; 30 µg), imipenem (IPM; 10 µg), meropenem (MEM; 10 µg), ceftazolin (CZ; 30 µg), ceftizoxime (ZOX; 30 µg), cefuroxime (CXM; 30 µg), cefepime (FEP; 30 µg), chloramphenicol (C; 30 µg), nitrofurantoin (F/M; 300 µg), nalidixic acid (NA; 30 µg), tetracycline (TE; 30 µg) and trimethoprim–sulphamethoxazole (SXT; 1.25 and 23.75 µg). Overnight nutrient broth cultures of the test strains were used and the turbidity of the inoculum as adjusted in phosphate-buffered saline (pH 7.4) to a 0.5 McFarland opacity standard (Becton and Dickinson). These cultures were then streaked onto Mueller–Hinton agar plates (Difco) using a sterile cotton swab. The antibiotic discs were dispensed using a disc dispenser (Becton and Dickinson) sufficiently separated from each other so as to avoid overlapping of inhibition zones. After 30 min, the plates were inverted and incubated at 37°C for 16–18 h. The isolates were considered sensitive according to the manufacturer's instructions.

Reference strains of *Escherichia coli* ATCC 25922 and *P. aeruginosa* ATCC 27853, as recommended by NCCLS (1997), were used as control organisms for verification of the anti-bacterial effect of the discs on Mueller–Hinton agar plates.

All discs were purchased from Becton and Dickinson.

For all isolates, we calculated the MAR index values (a/b , where a represents the number of antibiotics the isolate was resistant to and b represents the total number of antibiotics the isolate was tested against). A MAR index value >0.2 is observed when isolates are exposed to a high risk sources of human or animal contamination, where antibiotics use is common. In contrast, a MAR index value ≤ 0.2 is observed when antibiotics are seldom or never used (Krumperman 1983).

Determination of the MIC of heavy metals

The minimal inhibitory concentration (MIC) of each isolate was determined by the plate dilution method (Summers and Silver 1972). Six different heavy metals (Cd^{+2} , Cu^{+2} , Cr^{+3} , Pb^{+2} , Mn^{+2} and Zn^{+2}) were used as $\text{CdCl}_2 \cdot 2\text{H}_2\text{O}$, $\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$, CrCl_3 , $\text{Pb}(\text{NO}_3)_2$, $\text{MnCl}_2 \cdot 2\text{H}_2\text{O}$ and ZnCl_2 (Merck), each in ten concentrations ranging from 12.5 to $>3,200$ $\mu\text{g/ml}$ and added to Mueller–Hinton agar (Difco). The isolates were considered resistant if the MIC values exceeded that of the control organism. *E. coli* K-12 strain was used as the control organism as described by Malik and Jaiswal (2000) and Akinbowale et al. (2007).

Results and discussion

Aeromonas and *Pseudomonas* species and isolates

A total of 198 isolates were obtained, representing two *Aeromonas* species and their 60 isolates and six *Pseudomonas* species with their 138 isolates (Table 1). The frequency of the isolates varied markedly between sites, with most *Aeromonas* isolates recorded from site 1 and most of the *Pseudomonas* isolates recorded from sites 2 and 3 (Table 1).

Species characterisation revealed that amongst the *Aeromonas* spp., *A. hydrophila* was the dominant species (95.0%) amongst three sampling sites accounting for 57 of the 60 *Aeromonas*

Table 1 Numbers of *Aeromonas* spp. and *Pseudomonas* spp. isolated from three sites in Iskenderun Bay, Turkey (northeast Mediterranean Sea)

Species	Sources			Total
	Site 1	Site 2	Site 3	
<i>Aeromonas</i>				
<i>A. hydrophila</i>	39	7	11	57
<i>A. caviae</i>	–	–	3	3
<i>Pseudomonas</i>				
<i>P. aeruginosa</i>	8	4	16	28
<i>P. cepacia</i>	–	1	–	1
<i>P. diminuta</i>	–	1	–	1
<i>P. fluorescens</i>	6	25	3	34
<i>P. putida</i>	–	27	3	30
<i>P. stutzeri</i>	11	4	29	44
Total	64	69	65	198

isolates; only three isolates of *A. caviae* were found. This agrees with the results of Havelaar et al. (1990) from The Netherlands, which reported that *A. hydrophila* was the most frequently isolated species from distribution waters, whereas *A. caviae* and *A. sobria* were predominant in only a few water systems. *A. hydrophila* is a ubiquitous and opportunistic pathogen, which is part of the normal microbial flora of aquatic animals including fish, and *A. hydrophila* strains producing cytotoxins, proteases and aerolysin were commonly isolated from both healthy and moribund fish (Cahill 1990). Sewage and effluents from domestic water purification plants have been suspected as vehicles for transmission of pathogenic aeromonads in a number of clinical cases (Monfort and Baleux 1991).

In our study, amongst the *Pseudomonas* isolates, *Pseudomonas stutzeri* was the most prevalent species (31.9%), followed by *Pseudomonas fluorescens* (24.6%), *Pseudomonas putida* (21.7%) and *P. aeruginosa* (20.3%) (Table 1); only single isolates of *P. cepacia* and *P. diminuta*, were recorded.

Pseudomonas stutzeri is distributed widely in the environment, occupying diverse ecological niches, and has also been isolated as an opportunistic pathogen from humans (Lalucat et al. 2006). The second most prevalent *Pseudomonas*, *P. fluorescens*, is an emerging pathogen closely related to *P. aeruginosa*. *P. fluorescens* produces

a cholin-esterase (Roche et al. 1998), and some strains possess a high affinity γ -aminobutyric acid binding protein (Guthrie et al. 2000) that could adversely affect neurotransmission.

The third most prevalent *Pseudomonas*, *P. putida*, is a ubiquitous aerobic Gram-negative bacillus. It can be found in aquatic environments, in soils, on vegetation and even on some animals. It is widespread in the domestic and hospital environments, particularly in sites related to water and plumbing, and has been reported from patients with cystic fibrosis.

Kueh et al. (1992) investigated the possibility of wound infection with sewage-related organisms following exposure to contaminated sea water. These authors found that *A. hydrophila*, *Vibrio parahaemolyticus*, *Vibrio alginolyticus* and *Vibrio cholerae* were amongst the bacteria able to cause different kind of infections, and they further observed that *Vibrionaceae* were present in coastal waters, even at unpolluted sites.

Antibiotic resistance patterns

In Iskenderun Bay, a high proportion of *Aeromonas* and *Pseudomonas* isolates were resistant to different anti-bacterial agents, although resistance to aminoglycosides, carbapenems and fourth-generation cephalosporins was relatively infrequent. *Aeromonas* isolates showed a high percentage resistance to cefazolin (66.6%) and trimethoprim–sulphamethoxazole (66.6%). A low percentage of *Aeromonas* isolates were resistant to amikacin (3.3%), gentamicin (13.3%), chloramphenicol (13.3%) and nalidixic acid (13.3%; Table 2). Ko et al. (1996) found that a high proportion of *Aeromonas* were resistant to trimethoprim–sulfamethoxazole and tetracycline.

There were some marked differences in the degree of antibiotic resistance between sites (Table 2). For *Aeromonas* strains, all from site 1 were susceptible to amikacin, gentamicin and chloramphenicol; all from site 2 were susceptible to amikacin, imipenem and tetracycline, whereas those from site 3 showed variable resistance to the antibiotics tested in this study. *Aeromonas* isolates from site 3 showed higher resistance to cefazolin (85.7%) than did those from site 1 (59%) and site

2 (71.4%); this may be because site 3 is nearer the settlement and industry area than are zones 1 and 2 (Fig. 1).

Recent studies have demonstrated that the presence of *Aeromonas* spp. in drinking water is a potential risk, since some species can produce a wide range of virulence factors (Warburton et al. 1994; Kühn et al. 1997; Brandi et al. 1999). A further concern is the increasing incidence of multi-drug resistance amongst *Aeromonas* spp. isolates that has been observed worldwide (Petersen and Dalsgaard 2003; Ottoviani et al. 2006; Matyar et al. 2007). Besides their potential pathogenicity to humans, many *Aeromonas* species have also been described as important fish pathogens (Hänninen et al. 1997; Doukas et al. 1998).

Amongst the *Pseudomonas* isolates in our study, a high percentage were resistant to nitrofurantoin (86.2%), cefazolin (84.8%) and cefuroxime (71.7%), and a low percentage were resistant to kanamycin (5.8%), meropenem (7.2%) cefepime (8.7%), amikacin (12.3%) and imipenem (16.7%). Resistance to imipenem was similar to the results of Blandino et al. (2004). Susceptibility to aminoglycosides and carbapenems was reported previously by Sader and Jones (2005) and is similar to our results. From the site 1, none of the *Pseudomonas* isolates was found resistant to amikacin.

Multiple antibiotic resistance index

The MAR index values ranged from 0.2 to 0.60 for *Aeromonas* strains and from 0.2 to 0.73 for the *Pseudomonas* strains (Fig. 2). A significant proportion of *Aeromonas* strains isolated from site 1 were resistant to seven antibiotics (30.8%), a significant proportion of *Pseudomonas* strains from site 2 were resistant to five antibiotics (42.5%) and a significant proportion of the *Pseudomonas* strains isolated from site 3 were resistant to six antibiotics (31.4%). A variety of profiles, involving resistance to tetracycline and chloramphenicol and other antibiotics, have been described by Radu et al. (2003), who detected multiple resistance of *Aeromonas* spp.

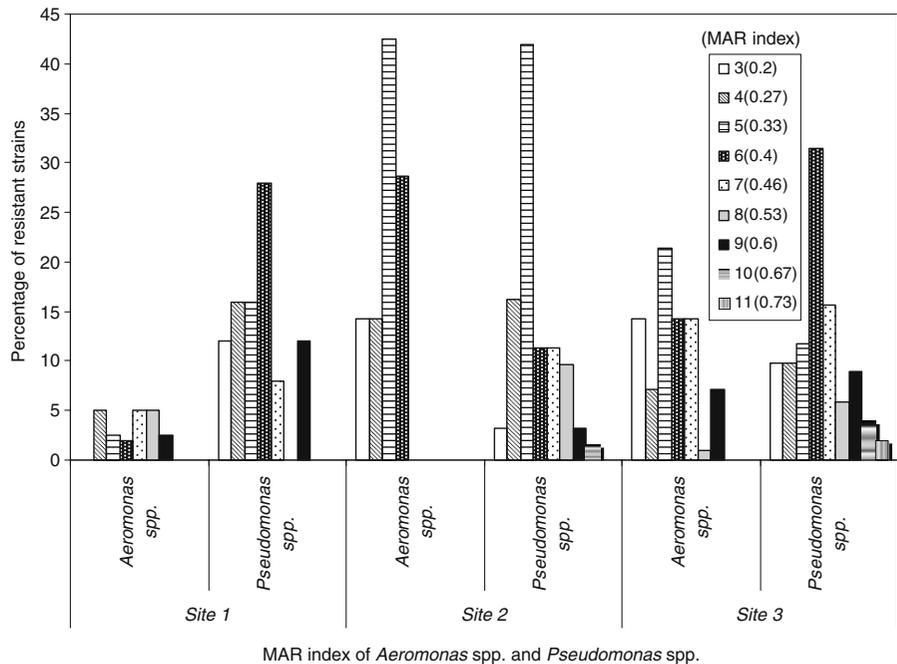
There was a similar level of resistance to nine antibiotics in *Aeromonas* strains (3.3%)

Table 2 Percentage of resistant isolates of *Aeromonas* spp. and *Pseudomonas* spp. from three sites in Iskenderun Bay, Turkey (northeast Mediterranean Sea) to 15 antibiotics belonging to the eight classes

Classes of antibiotics	Antibiotics	Site 1		Site 2		Site 3		Total	
		<i>Aeromonas</i> spp. (N = 39)	<i>Pseudomonas</i> spp. (N = 25)	<i>Aeromonas</i> spp. (N = 7)	<i>Pseudomonas</i> spp. (N = 62)	<i>Aeromonas</i> spp. (N = 14)	<i>Pseudomonas</i> spp. (N = 51)	<i>Aeromonas</i> spp. (N = 60)	<i>Pseudomonas</i> spp. (N = 138)
Percentage of resistant isolates									
Aminoglycosides	AN (30 µg)	-	-	-	6.4	14.3	23.5	3.3	12.3
	S (10 µg)	2.5	12	42.8	40.3	35.7	43.1	15.0	36.2
	GM (10 µg)	-	8	42.8	12.9	35.7	33.3	13.3	19.6
Carbapenems	K (30 µg)	23	16	14.2	1.6	14.2	5.8	20.0	5.8
	IPM (10 µg)	56.4	16	-	8	14.2	27.4	40.0	16.7
	MEM (10 µg)	51.2	12	14.2	4.8	21.4	7.8	40.0	7.2
Cephalosporins	CZ (30 µg)	59	68	71.4	90.3	85.7	86.2	66.6	84.8
	ZOX (30 µg)	61.5	28	14.2	1.6	35.7	17.6	50.0	12.3
	CXM (30 µg)	56.4	64	71.4	80.6	42.8	64.7	55.0	71.7
Chloramphenicol	FEP (30 µg)	53.8	28	42.8	3.2	14.2	5.8	43.3	8.7
	C (30 µg)	-	52	42.8	67.7	35.7	54.8	13.3	60.8
	F/M (300 µg)	5.1	80	71.4	95.1	50	78.4	23.3	86.2
Nitrofurantoin	NA (30 µg)	5.1	8	14.2	25.8	35.7	37.2	13.3	26.8
	TE (30 µg)	5.1	60	-	40.3	21.4	39.2	15.0	43.5
Trimethoprim-sulphamethoxazole	SXT (1.25 and 23.75 µg)	76.9	72	42.8	77.4	50	49	66.6	65.9

Number of isolates is shown at the top each column

Fig. 2 Anti-bacterial multi-resistance of *Aeromonas* spp. and *Pseudomonas* spp. isolated from three different sites in Iskenderun Bay, Turkey



and *Pseudomonas* strains (5.0%). A high multi-resistance incidence, similar to or higher than those found in the present study, has been reported in aquatic environments and fish isolates (Vivekanadhan et al. 2002; Matyar et al. 2007).

According to Anderson and Levin (1999), the frequency and rates of ascent and dissemination of antibiotic resistance in bacterial populations can be directly related to the applied compounds. De Vicente et al. (1990) reported that the number of strains multi-resistant to several antibiotics was high in seawater samples containing low levels of fecal indicators. A high incidence of antibiotic resistance in *Pseudomonas* spp. was reported by Jones et al. (1986) for lake waters.

Resistance to heavy metals

Trends in heavy metal resistance varied depending on the isolates for *Aeromonas*, Cu > Mn > Cr > Cd = Zn > Pb (Table 3) and for *Pseudomonas*, Cu > Cd > Mn > Zn > Cr > Pb (Table 4). Other studies have shown the following order of resistance for *Aeromonas* isolates, Cd > Cu > Hg > Cr (Miranda and Castillo 1998) and for *Pseudomonas* and *Micrococcus* isolates Mn > Cu > Pb > Zn > (Benka-Coker and Ekundayo

1998). In the present study in Iskenderun Bay, resistance to six heavy metals was as follows: for the *Aeromonas* and *Pseudomonas* isolates to cadmium, 35.0% and 56.5%; copper, 98.3% and 75.4%; chromium, 38.3% and 31.9%; lead, 1.7% and 7.2%; manganese, 43.3% and 44.9% and zinc 35.0% and 41.3%, respectively (Tables 3 and 4).

The MICs of the isolates ranged from 25 to >3,200 µg/ml. The *Aeromonas* isolates showed a higher resistance to copper and chromium than did the *Pseudomonas* isolates, and the *Pseudomonas* isolates showed a higher resistance to cadmium, lead, manganese and zinc than did the *Aeromonas* isolates. Tolerance to the maximum MIC (>3,200 µg/ml) for copper was 2.1% for *Aeromonas* and 1.6% for *Pseudomonas* isolates. *Aeromonas* strains from site 3 showed higher resistance to chromium, lead and manganese than did the *Aeromonas* strains from sites 1 and 2. Similarly, the *Pseudomonas* strains from site 3 showed higher resistance to chromium, lead, manganese and zinc than did *Pseudomonas* strains from sites 1 and 2. A possible explanation for these differences is that site 3 is nearer to the iron-steel factories than are sites 1 and 2 (Fig. 1). Higher proportion of metal resistance detected in the *Aeromonas* and *Pseudomonas* strains from site 3

Table 3 Heavy metal tolerance in *Aeromonas* spp. from three sites in Iskenderun Bay, Turkey (northeast Mediterranean Sea)

Heavy metal	Site of isolates	Number of total isolates	Metal concentrations (µg/ml) with number of tolerant isolates									Resistant isolates			
			12.5	25	50	100	200	400	800	1,600	3,200	>3,200	n	%	
Cadmium															
<i>Aeromonas</i>	Site 1	39				a								13	33.3
<i>Aeromonas</i>	Site 2	7			11	15	2	11						6	85.7
<i>Aeromonas</i>	Site 3	14		2	9	1	2	4	1	1				2	14.3
Total		60		2	20	17	4	15	1	1				21	35.0
Copper															
<i>Aeromonas</i>	Site 1	39						a						39	100.0
<i>Aeromonas</i>	Site 2	7					1		1			5		6	85.7
<i>Aeromonas</i>	Site 3	14								1	12	1		14	100.0
Total		60					1	39	1	1	17	1		59	98.3
Chromium															
<i>Aeromonas</i>	Site 1	39								a				10	25.6
<i>Aeromonas</i>	Site 2	7					1	4	2					–	–
<i>Aeromonas</i>	Site 3	14							1	2	11			13	92.8
Total		60					1	4	32	12	11			23	38.3
Lead															
<i>Aeromonas</i>	Site 1	39								a				–	–
<i>Aeromonas</i>	Site 2	7								12	27			–	–
<i>Aeromonas</i>	Site 3	14							1	8	4	1		1	7.1
Total		60							1	24	34	1		1	1.7
Manganese															
<i>Aeromonas</i>	Site 1	39								a				14	35.9
<i>Aeromonas</i>	Site 2	7								7	14	4	14	2	14.3
<i>Aeromonas</i>	Site 3	14							1	2	1	10		10	71.4
Total		60							1	9	18	6	26	26	43.3
Zinc															
<i>Aeromonas</i>	Site 1	39								a				18	46.1
<i>Aeromonas</i>	Site 2	7								12	9	3	15	3	42.8
<i>Aeromonas</i>	Site 3	14								4	7			–	–
Total		60								1	22	16	5	15	35.0

a minimal inhibition concentration of standard strain *E. coli* K12, n total number of tolerant isolates

could be the result of heavy metal contamination from iron-steel factories. Both Gram-positive and Gram-negative bacteria can resist heavy metals (Silver and Walderhaug 1992). Resistance of toxic metals in bacteria probably reflects the degree of environmental contamination with these substances and may be directly related to exposure of bacteria to them (Aiking et al. 1984).

Natural ecosystems containing high concentrations of heavy metals are also frequent. Not surprisingly, heavy metal resistance genes are commonly found in environmental bacteria (Silver and Phung 1996).

The present study is one of only a few that addresses the incidence of antibiotic and heavy metal resistance in *Aeromonas* spp. and *Pseudomonas* spp. With all six of the metals tested, there was a high frequency of cefazolin resistance amongst all the isolates.

The *Aeromonas* strains which were metal resistant also showed a high resistance to antibiotics as follows: from site 1, high resistance to cefazolin, cefuroxime and trimethoprim-sulphamethoxazole; from site 2, high resistance to cefazolin and cefuroxime and from site 3, high resistance to cefazolin. The *Pseudomonas*

Table 4 Heavy metal tolerance in *Pseudomonas* spp. from three sites in Iskenderun Bay, Turkey (northeast Mediterranean Sea)

Heavy metal	Site of isolates	Number of total isolates	Metal concentrations (µg/ml) with number of tolerant isolates									Resistant isolates		
			12.5	25	50	100	200	400	800	1,600	3,200	>3,200	n	%
Cadmium			a											
<i>Pseudomonas</i>	Site 1	25		1	11	8	3	2					5	20.0
<i>Pseudomonas</i>	Site 2	62		3	8	8	11	22	5	5			43	69.3
<i>Pseudomonas</i>	Site 3	51		2	12	7	4	4	19	2	1		30	58.8
Total		138		6	31	23	18	28	24	7	1		78	56.5
Copper			a											
<i>Pseudomonas</i>	Site 1	25					2	17			6		23	92.0
<i>Pseudomonas</i>	Site 2	62					8	11	7	9	27		54	87.1
<i>Pseudomonas</i>	Site 3	51			3	21	1	9			14	3	27	52.9
Total		138			3	31	29	16	9	47	3		104	75.4
Chromium			a											
<i>Pseudomonas</i>	Site 1	25						13	12				–	–
<i>Pseudomonas</i>	Site 2	62			1	1	22	32	3	3			6	24.0
<i>Pseudomonas</i>	Site 3	51						2	11	20	18		38	74.5
Total		138			1	1	37	55	23	21	3		44	31.9
Lead			a											
<i>Pseudomonas</i>	Site 1	25			1	3	13		8				–	–
<i>Pseudomonas</i>	Site 2	62			1		23	15	20	3			3	4.8
<i>Pseudomonas</i>	Site 3	51				1	7	18	18	7			7	13.7
Total		138			2	4	43	33	46	10			10	7.2
Manganese			a											
<i>Pseudomonas</i>	Site 1	25				7	11	4	1	2			2	8.0
<i>Pseudomonas</i>	Site 2	62				1	8	14	10	29			29	46.8
<i>Pseudomonas</i>	Site 3	51					9	7	4	29	2		31	60.8
Total		138				8	28	25	15	60	2		62	44.9
Zinc			a											
<i>Pseudomonas</i>	Site 1	25			2	19	3	1					4	16.0
<i>Pseudomonas</i>	Site 2	62			7	34	10		4	7			21	33.9
<i>Pseudomonas</i>	Site 3	51			10	9	5	3	21	2	1		32	62.7
Total		138			2	17	62	18	4	25	9	1	57	41.3

a minimal inhibition concentration of standard strain *E. coli* K12, n total number of tolerant isolates

strains which were metal resistant also showed a high resistance to antibiotics as follows: from site 1, high resistance to cefazolin, cefuroxime, nitrofurantoin, tetracycline and trimethoprim-sulphamethoxazole; from site 2, showed a high resistance to five antibiotics: cefazolin, cefuroxime, chloramphenicol, nitrofurantoin and trimethoprim-sulphamethoxazole and from site 3, high resistance to four antibiotics: cefazolin, cefuroxime, chloramphenicol and nitrofurantoin.

Industrial activities, mining and intensive farming are causing dramatic changes in natural ecosystems. Amongst the novel selective pres-

ures that face environmental bacterial populations are discharges of heavy metals, xenobiotic compounds, antibiotics and organic solvents, all of which can have a significant impact on the environmental selection of antibiotic resistance genes (Alonso et al. 2001). Some of these microorganisms that possess resistance genes could cause important diseases.

Conclusion and recommendations

The presence, antibiotic resistance and heavy metal resistance patterns of *Aeromonas* strains

and *Pseudomonas* strains in a marine environment at northeast of Mediterranean Sea were investigated. These results suggests that Iskenderun Bay, Turkey, is a reservoir for antibiotic and metal resistant *Aeromonas* and *Pseudomonas* strains and might be increasing the numbers of resistant bacteria in the Bay with possibilities of transferring of their resistance determinants to other pathogenic bacteria. Although Chang and Bolton (1987) found that plasmid-mediated antibiotic resistance in *Aeromonas* spp. was not frequent, McIntosh et al. (2008) have found an important incidence of R-plasmids in multiple antibiotic and mercury resistant *Aeromonas* isolates from Atlantic salmon. The study of Rhodes et al. (2000) provided evidence that related tetracycline-resistance encoding plasmids have been transferred between different *Aeromonas* species and *E. coli* and between the hospital and aquaculture environments in distinct geographical locations.

Thus, plasmid-mediated antibiotic and heavy metal resistance in *Aeromonas* spp. in the Mediterranean Sea should be further elucidated. Further works are needed to understand better metal and antibiotic resistant *Aeromonas* spp. and *Pseudomonas* spp. along with other parts of the Turkish coasts on the Mediterranean and Aegean Seas.

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