

## Resistances in *Aeromonas* Species from the SENTRY Global Surveillance

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### Introduction

*Aeromonas* spp. are uncommon pathogens often causing severe illness. Although they harbour specific  $\beta$ -lactamases, resistance to many other drugs should be low as their reservoir is environmental. We examined the resistance patterns of all *Aeromonas* spp. isolated in the SENTRY surveillance program over 5 years (1997 - 2001).

### Methods

#### Isolates

All *Aeromonas* isolated from blood, lower respiratory tract, skin/soft tissue, urine and feces. A total of 57 laboratories from 24 countries contributed strains.

#### Susceptibility testing

All isolates were tested by broth microdilution using custom made panels (TREK™ Diagnostic Systems), according to NCCLS standards,<sup>1</sup> to over 25 antimicrobials. Strains were defined as resistant if the MIC exceeded NCCLS *Enterobacteriaceae* breakpoints for "susceptible"<sup>2</sup>

Multi-resistant *Aeromonas* spp. were defined as those strains demonstrating resistance to 4 or more of the following agents: cefoxitin, ceftriaxone, ceftazidime, cefepime, ciprofloxacin, gentamicin, tetracycline, meropenem or trimethoprim/sulphamethoxazole.

### Results

- A total of 258 strains were collected during the study period. Blood isolates accounted for 50% of all isolates (Figure 1)
- Over 87% of all species were either *Aeromonas hydrophila* (n=149), *A. caviae* (n=40) or *A. sobria* (n=37) (Figure 2)
- Resistance rates for the different species and regions are shown in Table 1
- Significant variation in resistance patterns were noted among species and by region
- Nalidixic acid resistance was high (49%) for *A. sobria* however no ciprofloxacin resistance was detected
- Multi-drug resistant strains were common (17%) in Latin American isolates (Figure 3). One strain of *S. salmonicida* from Venezuela was resistant to all drugs tested. It had a cefepime MIC of 16 mg/L
- Only 2% of isolates were meropenem resistant
- Ceftriaxone-resistant strains were resistant to ceftazidime and had elevated MICs to cefepime and aztreonam. Their meropenem MICs were not elevated

Figure 1.  
Source of Isolates

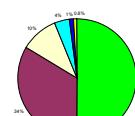


Figure 2.  
Species Distribution by Region

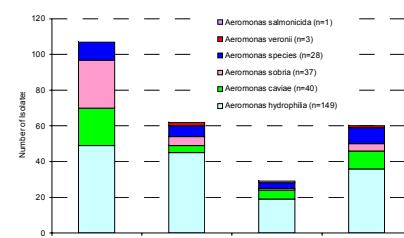
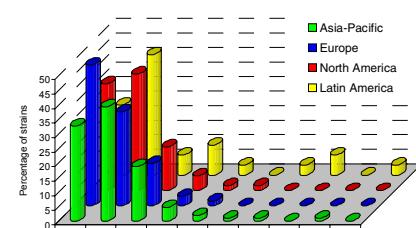


Figure 3.  
Number of Resistances by Region



#### Results (Continued)

- There was a statistical association between tetracycline and trimethoprim/sulpha resistance (Table 3), and between ciprofloxacin and gentamicin resistance (Table 4)

Table 1. Resistance in *Aeromonas* spp. - a Global Perspective

Antimicrobial Agent	MIC (mg/L)	Number (%) of isolates resistant by species						Number (%) of isolates resistant by region			
		<i>A. hydrophilia</i> (n=149)	<i>A. caviae</i> (n=40)	<i>A. sobria</i> (n=37)	<i>Aeromonas</i> sp. (n=28)	<i>A. veronii</i> (n=3)	Asia-Pacific (n=107)	Europe (n=62)	Latin America (n=29)	N. America (n=60)	Global (n=258)
Amoxicillin/clavulanate	>8	137 (91.9)	31 (77.5)	31 (83.8)	28 (100)	2	95 (88.8)	53 (85.5)	28 (96.6)	54 (90.0)	230 (89.1)
Ticarcillin/clavulanate	>16	100 (67.1)	24 (60.0)	26 (70.3)	22 (78.6)	1	64 (59.8)	39 (62.9)	26 (89.7)	45 (75.0)	174 (67.4)
Piperacillin/tazobactam	>16	39 (26.2)	8 (20.0)	14 (37.8)	6 (21.4)	1	22 (20.6)	23 (37.1)	8 (27.6)	16 (26.7)	69 (26.7)
Cefoxitin	>8	60 (40.3)	16 (40.0)	6 (16.2)	15 (53.6)	0	46 (43.0)	18 (29.0)	13 (44.8)	21 (35.0)	98 (38.0)
Ceftriaxone	>8	7 (4.7)	6 (15.0)	0	4 (14.3)	0	5 (4.7)	2 (3.2)	6 (20.7)	5 (8.3)	18 (7.0)
Ceftazidime	>8	5 (3.4)	4 (10.0)	0	4 (14.3)	0	5 (4.7)	1 (1.6)	4 (13.8)	4 (6.7)	14 (5.4)
Cefepime	>8	0	2 (5.0)	0	0	0	1 (0.9)	0	2 (6.9)	0	3 (1.2)
Aztreonam	>8	2 (1.3)	2 (5.0)	0	1 (3.6)	0	1 (0.9)	1 (1.6)	3 (10.3)	1 (1.7)	6 (2.3)
Meropenem	>4	0	0	2 (5.4)	3 (10.7)	0	4 (3.7)	0	1 (3.4)	1 (1.7)	6 (2.3)
Gentamicin	>4	3 (2.0)	1 (2.5)	2 (5.4)	2 (7.1)	0	3 (2.8)	1 (1.6)	3 (10.3)	2 (3.3)	9 (3.5)
Nalidixic acid	>16	36 (24.2)	7 (17.5)	18 (48.6)	6 (21.4)	1	31 (29.0)	22 (35.5)	8 (27.6)	8 (13.3)	69 (26.7)
Ciprofloxacin	>1	2 (1.3)	2 (5.0)	0	1 (3.6)	0	1 (0.9)	1 (1.6)	3 (10.3)	1 (1.7)	6 (2.3)
Tetracycline	>4	29 (19.5)	9 (22.5)	17 (45.9)	8 (28.6)	1	39 (36.4)	12 (19.4)	10 (34.5)	4 (6.7)	65 (25.2)
Trimethoprim/sulpha	>1	23 (15.4)	3 (7.5)	9 (24.3)	6 (21.4)	1	20 (18.7)	8 (12.9)	10 (34.5)	5 (8.3)	43 (16.7)
Polymyxin B	>8	28 (35.0)	0	6 (23.1)	2 (15.4)	2	30 (38.5)	2 (5.1)	2 (20.0)	2 (9.5)	36 (24.3)

Table 2.  
Tetracycline vs Trimeth/sulphamethoxazole

Tetracycline				
	≤4	8	≥8	Total
Sulpha/ trimeth	≤1	180	8	215
	2	13	7	43
Total	193	7	50	258

Table 3. Ciprofloxacin vs Gentamicin

GEN	Ciprofloxacin			
	≤4	1	2	≥4
≤1	247	1	1	249
8	4	0	0	4
≥16	1	0	4	5
Total	252	1	5	258

### Conclusions

- Resistance patterns among *Aeromonas* species showed considerable variation
- Regional differences in resistance levels were also evident
- High frequencies of TET and SXT resistance suggest significant environmental selection pressure
- Possible extended-spectrum and metallo- $\beta$ -lactamases were noted

#### Acknowledgments

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#### References

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