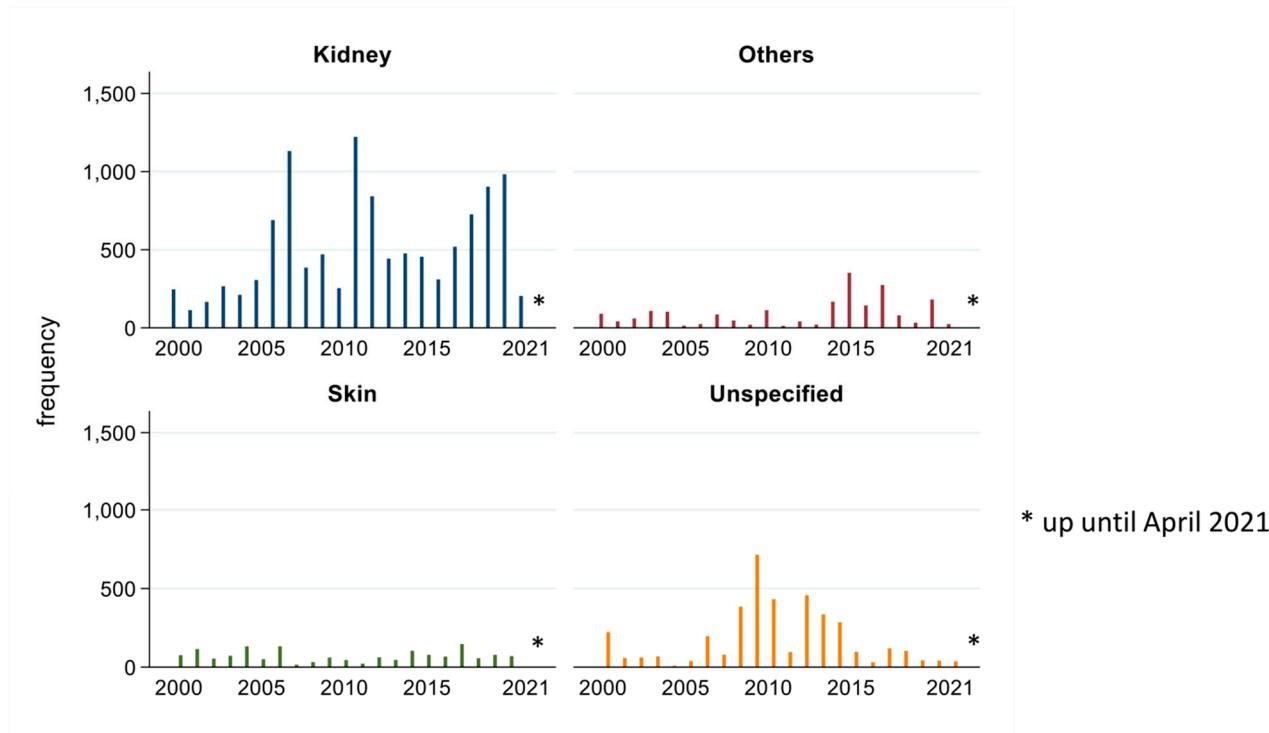
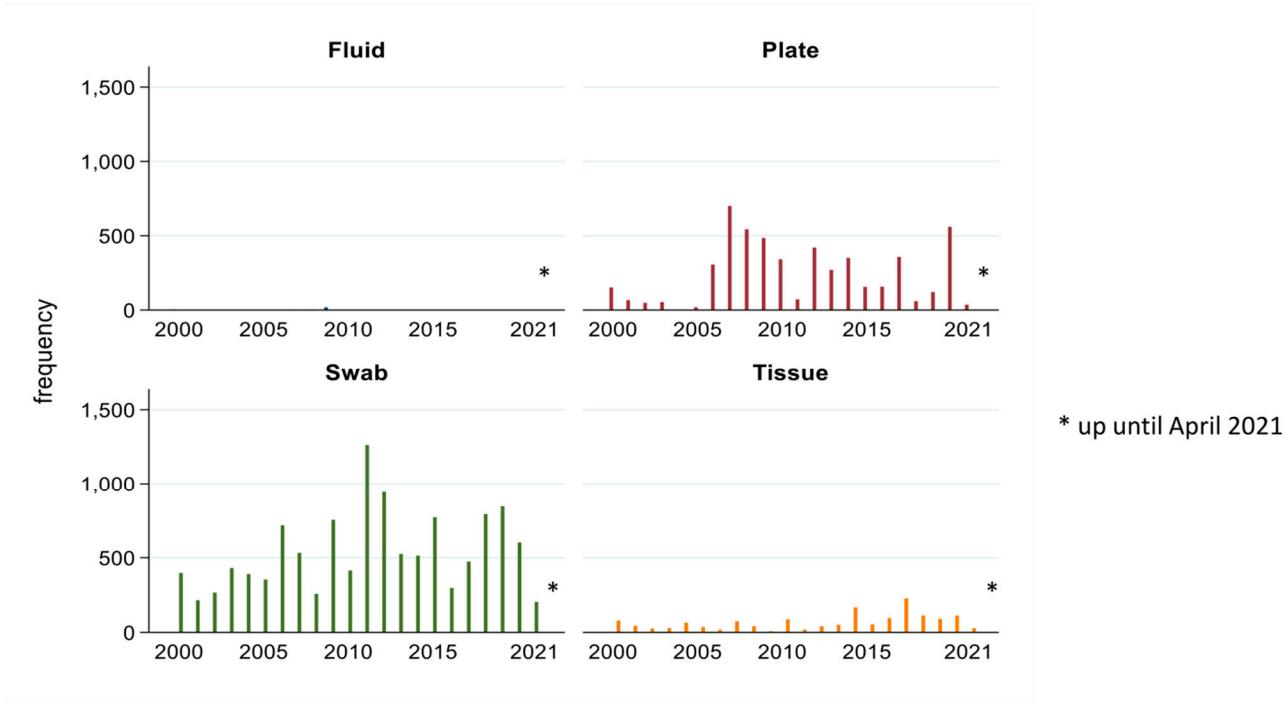


# Antimicrobial Susceptibility Profiles of Bacteria Commonly Isolated from Farmed Salmonids in Atlantic Canada (2000–2021)

Rasaq Abiola Ojasanya <sup>1,\*</sup>, Ian A. Gardner <sup>1</sup>, David B. Groman <sup>2</sup>, Sonja Saksida <sup>1</sup>, Matthew E. Saab <sup>2</sup> and Krishna Kumar Thakur <sup>1</sup>



**Figure S1.** Annual frequency of salmonid samples submitted for bacteriology to AVC ADSBL from 2000 to 2021 by anatomic site of sample collection.



**Figure S2.** Annual frequency of salmonid specimens submitted for bacteriology to AVC ADSBL from 2000 to 2021.

**Table S1.** Frequency of samples originating from different anatomic sites submitted to the AVC ADSBL (2000-2021).

Species	Kidney (n = 11,320)	Skin (n = 1517)	Unspecified (n = 3905)	Others (n = 2034)
n (%)				
Atlantic salmon	9994 (88.29)	1392 (91.76)	3700 (94.75)	1782 (87.61)
Rainbow trout	1326 (11.71)	125 (8.24)	205 (5.25)	252 (12.39)

**Table S2.** Bacterial isolates detected from different anatomic sites of salmonids samples submitted to the AVC ADSBL (2000-2021).

Bacterial Isolates	Kidney (n = 11,320)	Skin (n = 1517)	Unspecified (n = 3905)	Others (n = 2034)
	%			
Mixed microbiota	9.5	23.7	17.4	17.4
Gram-negative bacilli	5.4	7.0	16.8	16.8
<i>Vibrio</i> spp.	5.6	15.2	8.9	7.0
<i>Pseudomonas</i> spp.	2.4	2.8	3.2	4.9
<i>Aeromonas</i> spp.	2.2	2.9	2.4	3.2
<i>Flavobacterium</i> spp.	0.3	1.9	0.6	4.7
Gram-positive bacilli	1.9	0.7	2.4	2.4
<i>Yersinia</i> spp.	1.4	0	1.1	1.1
<i>Vibrio anguillarum</i>	0.1	0	1.2	1.2
<i>Mycobacterium</i> spp.	0.3	0.1	1.2	0.1
<i>Aeromonas sobria</i>	0.3	0.1	0.6	0.6
<i>Edwardsiella piscicida</i>	1.4	0	0	0
Gram-positive bacteria	0.1	0	0.5	0.5
<i>Arthrobacter</i> spp.	0.4	0	0.3	0.3
<i>Enterobacter</i> spp.	0	0.5	0.2	0.2
<i>Serratia</i> spp.	0.1	0	0.4	0.4
<i>Aliivibrio salmonicida</i>	0.5	0	0.2	0.2
<i>Micrococcus</i> spp.	0	0	0.3	0.3
<i>Shewanella putrefaciens</i>	0	0.5	0	0
<i>Moraxella</i> spp.	0	0.1	0.2	0.2
<i>Vibrio alginolyticus</i>	0	0	0.2	0.2
<i>Bacillus</i> spp.	0	0	0.3	0.3
<i>Staphylococcus</i> spp.	0.1	0	0.1	0.1
<i>Rhodococcus</i> spp.	0.1	0	0.1	0.1
<i>Lactobacillus</i> spp.	0.1	0	0.1	0.1
<i>Hafnia alvei</i>	0.2	0	0	0
<i>Plesiomonas</i> spp.	0.2	0	0	0

<i>Staphylococcus epidermidis</i>	0	0	0.1	0.1
<i>Streptococcus</i> spp.	0	0	0.1	0.1
<i>Yersinia ruckeri</i> type 2	0.2	0	0	0
<i>Aliivibrio fisherii</i>	0	0.1	0	0
<i>Carnobacterium maltaromaticum</i>	0.1	0	0	0
<i>Enterococcus</i> spp.	0.1	0	0	0
<i>Photobacterium</i> spp.	0.1	0	0	0
<i>Proteus mirabilis</i>	0.1	0	0	0
<i>Proteus</i> spp.	0.1	0	0	0
<i>Pseudomonas putida</i>	0.1	0	0	0
<i>Yersinia ruckeri</i>	0.1	0	0	0
<i>Corynebacterium</i> spp.	0	0.1	0	0

n = number of isolates, Unspecified- sample sites without clear identification. Others-include sample sites from the abdominal cavity, blood, bladder, brain, ova, eye, fin, gill, head, heart, intestine, jaw, liver, mouth, muscle, ovarian fluid, peritoneal cavity, spleen, tail, vent, wound, and yolk sac.

**Table S3.** Frequency of bacterial isolates detection in submitted cases to the AVC ADSBL by species of salmonid samples (2000-2021).

Bacterial Isolates	Frequency of Cases (n)	Atlantic Salmon	Rainbow Trout
		n (%)	n (%)
Mixed microbiota	300	281 (93.7)	19 (6.3)
Gram-negative bacilli	284	269 (94.7)	15 (5.3)
<i>Vibrio</i> spp	150	145 (96.7)	5 (3.3)
Gram-positive bacilli	54	47 (87.0)	7 (13.0)
<i>Aeromonas</i> spp.	35	34 (97.1)	1 (2.86)
<i>Mycobacterium</i> spp.	19	19 (100.0)	0
<i>Flavobacterium</i> spp.	14	11 (78.57)	3 (21.4)
<i>Yersinia</i> spp.	12	12 (100.0)	0
<i>Nocardia</i> spp.	13	13 (100.0)	0
Gram-positive cocci	9	6 (66.7)	3 (33.3)
Gram-negative bacteria	9	5 (55.6)	4 (44.4)
<i>Bacillus</i> spp.	5	5 (100.0)	0
Gram-positive bacteria	5	5 (100.0)	0
<i>Lactobacillus</i> spp.	4	4 (100.0)	0
<i>Rhodococcus</i> spp.	4	4 (100.0)	0

<i>Serratia liquefaciens</i>	4	4 (100.0)	0
<i>Staphylococcus</i> spp.	4	4 (100.0)	0
<i>Vibrio anguillarum</i> type 2	4	4 (100.0)	0
<i>Yersinia ruckeri</i> type 2	3	3 (100.0)	0
<i>Enterobacter</i> spp.	3	3 (100.0)	0
<i>Plesiomonas</i> spp.	3	3 (100.0)	0
<i>Carnobacterium maltaromaticum</i>	2	1 (50.0)	1 (50.0)
<i>Aliivibrio fischeri</i>	2	2 (100.0)	0
Gram-positive coccobacillus	2	2 (100.0)	0
<i>Micrococcus</i> spp.	2	2 (100.0)	0
<i>Acinetobacter</i> spp.	1	1 (100.0)	0
<i>Chryseobacterium</i> spp.	1	1 (100.0)	0
<i>Lactococcus lactis</i>	1	0	1 (100.0)
<i>Photobacterium damsel</i>	1	0	1 (100.0)
<i>Photobacterium</i> spp.	1	0	1 (100.0)
<i>Proteus vulgaris</i>	1	1 (100.0)	0
<i>Pseudomonas chlororaphis</i>	1	1 (100.0)	0
<i>Pseudomonas putida</i>	1	1 (100.0)	0
<i>Serratia</i> spp.	1	1 (100.0)	0
<i>Staphylococcus epidermidis</i>	1	1 (100.0)	0
<i>Streptococcus</i> spp.	1	1 (100.0)	0
<i>Corynebacterium</i> spp.	1	0	1 (100.0)
<i>Vibrio anguillarum</i>	1	1 (100.0)	0
<i>Vibrio fischeri</i>	1	1 (100.0)	0

**Table S4.** Antimicrobial susceptibility patterns (% S, I, and R) for tested antimicrobials in bacterial isolates from samples of salmonid. (2000-2021)

Bacterial Isolates	Florfenicol				Oxytetracycline				Ormetoprim-sulfadimethoxine				Trimethoprim-sulfamethoxazole				
	N	n	S	I	R	n	S	I	R	n	S	I	R	n	S	I	R
<i>Vibrio</i> spp.	141	140	100.0	0	0	141	91.0	2.1	7.1	63	45.0	0	0	138	97.2	0	0.7
<i>Pseudomonas</i> spp.	53	53	7.6	9.4	83.0	53	70.0	15.1	15.1	7	0	0	13.2	53	22.6	11.3	66.0
<i>Aeromonas</i> spp.	45	44	91.1	2.2	4.4	44	36.0	2.2	60.0	6	13.3	0	0	44	91.1	0	6.7
<i>Yersinia</i> spp.	28	27	100.0	0	0	28	96.0	0	3.6	25	3.9	0	0	28	100.0	0	0

<i>Flavobacterium</i> spp.	21	21	85.7	0	14.3	21	67.0	0	33.3	3	9.5	0	4.8	18	57.1	0	28.6
<i>Nocardia</i> spp.	12	12	8.3	42.0	50.0	12	8.3	25.0	66.7	5	41.7	0	0	12	100.0	0	0
<b>Enrofloxacin</b>										<b>Ampicillin</b>							
<b>Bacterial Isolates</b>	<b>N</b>	<b>n</b>	<b>S</b>	<b>I</b>	<b>R</b>	<b>n</b>	<b>S</b>	<b>I</b>	<b>R</b>								
<i>Vibrio</i> spp.	141	141	100.0	0	0	141	48.9	4.3	46.8								
<i>Pseudomonas</i> spp.	53	50	86.8	7.6	0	53	1.9	0	98.1								
<i>Aeromonas</i> spp.	45	41	91.1	0	0	44	6.7	0	91.1								
<i>Yersinia</i> spp.	28	28	100.0	0	0	28	64.3	3.6	32.1								
<i>Flavobacterium</i> spp.	21	21	100.0	0	0	21	23.8	4.8	71.4								
<i>Nocardia</i> spp.	12	12	100.0	0	0	12	0	0	100.0								

N= Total number of isolates, n=number of isolates that were tested for antimicrobial susceptibility, S- Susceptible, I- Intermediate, R- Resistant cumulative percentages may not add up to 100 as not all isolates had susceptibility test.