

# Phenotypic and genotypic characteristics of antimicrobial resistance in *Citrobacter freundii* isolated from domestic ducks (*Anas platyrhynchos domesticus*) in Bangladesh

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**Supplementary Table S1:** Pearson correlation coefficients assessing correlation between pairs of antibiotics to which *C. freundii* isolates showed resistance.

		C	CN	CIP	CL	AZM	TE	AMP	CRO	COT	FO	F	CTX	CAZ	LEV
<b>C</b>	ρ	.a	.a												
	Sig. (2-tailed)		.												
<b>CN</b>	ρ	.a	1												
	Sig. (2-tailed)	.	.												
<b>CIP</b>	ρ	.a	<b>.749**</b>	1											
	Sig. (2-tailed)	.	<b>0.000</b>	.											
<b>CL</b>	ρ	.a	0.115	0.153	1										
	Sig. (2-tailed)	.	0.585	0.465	.										
<b>AZM</b>	ρ	.a	<b>.498*</b>	<b>.665**</b>	0.23	1									
	Sig. (2-tailed)	.	<b>0.011</b>	<b>0.000</b>	0.268	.									
<b>TE</b>	ρ	.a	<b>.634**</b>	<b>.846**</b>	0.181	<b>.786**</b>	1								
	Sig. (2-tailed)	.	<b>0.001</b>	<b>0.000</b>	0.387	<b>0.000</b>	.								
<b>AMP</b>	ρ	.a	0.316	<b>.421*</b>	-0.115	<b>.445*</b>	0.309	1							
	Sig. (2-tailed)	.	0.124	<b>0.036</b>	0.585	<b>0.026</b>	0.132	.							
<b>CRO</b>	ρ	.a	0.081	0.236	0.075	0.327	<b>.417*</b>	0.208	1						
	Sig. (2-tailed)	.	0.701	0.256	0.72	0.11	<b>0.038</b>	0.32	.						
<b>COT</b>	ρ	.a	<b>.688**</b>	<b>.919**</b>	0.167	<b>.724**</b>	<b>.921**</b>	<b>.459*</b>	<b>.452*</b>	1					
	Sig. (2-tailed)	.	<b>0.000</b>	<b>0.000</b>	0.426	<b>0.000</b>	<b>0.000</b>	<b>0.021</b>	<b>0.023</b>	.					
<b>FO</b>	ρ	.a	-0.115	-0.153	0.042	0.181	-0.181	0.115	-0.075	-0.167	1				
	Sig. (2-tailed)	.	0.585	0.465	0.843	0.387	0.387	0.585	0.72	0.426	.				
<b>F</b>	ρ	.a	.a	.a	.a	.a	.a	.a	.a	.a	.a	.a			
	Sig. (2-tailed)	.	.	.	.	.	.	.	.	.	.	.			
<b>CTX</b>	ρ	.a	0.187	0.25	0.102	0.04	0.161	0.281	<b>.431*</b>	0.204	-0.102	.a	1		
	Sig. (2-tailed)	.	0.37	0.228	0.627	0.848	0.442	0.174	<b>0.032</b>	0.328	0.627	.	.		
<b>CAZ</b>	ρ	.a	0.081	0.236	0.075	0.079	0.169	0.208	<b>.621**</b>	0.201	-0.075	.a	<b>.739**</b>	1	
	Sig. (2-tailed)	.	0.701	0.256	0.72	0.706	0.42	0.32	<b>0.001</b>	0.335	0.72	.	<b>0.000</b>	.	

LEV	q	.a	.554**	.653**	0.153	.497*	.678**	0.226	0.236	.578**	-0.153	.a	0.25	.492*	1
	Sig. (2-tailed)	.	0.004	0.000	0.465	0.012	0.000	0.277	0.256	0.002	0.465	.	0.228	0.012	.

Here, \*\* Correlation is significant at the 0.01 level (2-tailed); \* Correlation is significant at the 0.05 level (2-tailed); .aCannot be computed because at least one of the variables is constant; q = Pearson correlation coefficient; LEV = levofloxacin; CAZ = ceftazidime; CTX = cefotaxime; F = nitrofurantoin; FO = fosfomycin; COT = cotrimoxazole; CRO = ceftriaxone; AMP = ampicillin; TE = tetracycline; AZM = azithromycin; CL = Cephalexin; CIP = ciprofloxacin; CN = gentamycin; C = chloramphenicol.

**Supplementary Table S2.** Pearson correlation coefficients assessing correlation between pairs of antibiotic resistance genes in *C. freundii* isolates from cloacal swabs of ducks.

		<i>bla</i> <sub>TEM-1</sub>	<i>bla</i> <sub>CMY-2</sub>	<i>bla</i> <sub>CMY-9</sub>	<i>bla</i> <sub>CTX-M-1</sub>	<i>bla</i> <sub>CTX-M-2</sub>	<i>bla</i> <sub>CTX-M-14</sub>	<i>bla</i> <sub>SHV-1</sub>	<i>sul1</i>	<i>sul2</i>	<i>tetA</i>	<i>tetB</i>	<i>tetC</i>	<i>qnrA</i>	<i>qnrB</i>	<i>qnrS</i>	<i>aacc2</i>	<i>aacc4</i>
<i>bla</i> <sub>TEM-1</sub>	Q	1																
	<i>p</i>																	
<i>bla</i> <sub>CMY-2</sub>	Q	<b>.417*</b>	1															
	<i>p</i>	<b>0.038</b>																
<i>bla</i> <sub>CMY-9</sub>	Q	0.109	0.261	1														
	<i>p</i>	0.604	0.207															
<i>bla</i> <sub>CTX-M-1</sub>	Q	.b	.b	.b	.b													
	<i>p</i>	.	.	.														
<i>bla</i> <sub>CTX-M-2</sub>	Q	.b	.b	.b	.b	.b												
	<i>p</i>	.	.	.	.													
<i>bla</i> <sub>CTX-M-14</sub>	Q	0.185	<b>.443*</b>	-0.147	.b	.b	1											
	<i>p</i>	0.377	<b>0.026</b>	0.482	.	.												
<i>bla</i> <sub>SHV-1</sub>	Q	.b	.b	.b	.b	.b	.b	.b										
	<i>p</i>	.	.	.	.	.	.											
<i>sul1</i>	Q	0.384	0.277	-0.012	.b	.b	0.08	.b	1									
	<i>p</i>	0.058	0.179	0.955	.	.	0.704	.										
<i>sul2</i>	Q	0.208	0.309	-0.166	.b	.b	<b>.421*</b>	.b	<b>.540**</b>	1								
	<i>p</i>	0.32	0.132	0.429	.	.	<b>0.036</b>	.	<b>0.005</b>									
<i>tetA</i>	Q	0.253	0.263	-0.202	.b	.b	0.086	.b	<b>.487*</b>	0.217	1							
	<i>p</i>	0.222	0.205	0.332	.	.	0.684	.	<b>0.013</b>	0.298								
<i>tetB</i>	Q	0.075	0.181	-0.06	.b	.b	<b>.408*</b>	.b	0.196	0.363	-0.14	1						
	<i>p</i>	0.72	0.387	0.775	.	.	<b>0.043</b>	.	0.347	0.074	0.504							
<i>tetC</i>	Q	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b					
	<i>p</i>	.	.	.	.	.	.	.	.	.	.	.	.					

qnrA	q	0.075	0.181	-0.06	.b	.b	.408*	.b	-0.212	-0.115	0.298	-0.042	.b	1				
	p	0.72	0.387	0.775	.	.	0.043	.	0.308	0.585	0.149	0.843	.					
qnrB	q	-0.242	0.079	-0.109	.b	.b	0.123	.b	-0.138	0.081	0.274	-0.075	.b	.553**	1			
	p	0.243	0.706	0.604	.	.	0.558	.	0.511	0.701	0.184	0.72	.	0.004				
qnrS	q	0.075	0.181	-0.06	.b	.b	-0.102	.b	0.196	-0.115	0.298	-0.042	.b	-0.042	-0.075	1		
	p	0.72	0.387	0.775	.	.	0.627	.	0.347	0.585	0.149	0.843	.	0.843	0.72			
aacc2	q	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	.b	
	p	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	.	
aacc4	q	0.161	-0.053	-0.129	.b	.b	-0.218	.b	.419*	0.01	.636**	-0.089	.b	-0.089	-0.161	.468*	.b	1
	p	0.442	0.802	0.54	.	.	0.295	.	0.037	0.961	0.001	0.672	.	0.672	0.442	0.018	.	

Here, \*\*Correlation is significant at the 0.01 level (2-tailed); \*Correlation is significant at the 0.05 level (2-tailed); .bCannot be computed because at least one of the variables is constant; q = Pearson correlation coefficient; p = Significance (2-tailed)