Mossambicus tilapia (*Oreochromis mossambicus*) collected from water bodies impacted by urban waste carries extended-spectrum beta-lactamases and integron-bearing gut bacteria

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Supplementary material

Supplementary table 1. Prevalence of resistance factors in the bacterial strains isolated from Tilapia gut

Isolate code	Identification	*IntI1*	*IntI2*	bla_{SHV}	bla_{OXA}	*aac(6')-Ib-cr*	bla_{CTX-M}
TAL_108	*Acinetobacter johnsonii*	+	+			+	
TAL_117	*Acinetobacter johnsonii*	+				+	+
TAL_128	*Acinetobacter johnsonii*			+			
WAK_92	*Aeromonas* sp.		+				
AUN_36	*Aeromonas aquariorum*						
AUN_39	*Aeromonas aquariorum*		+				
TAL_112	*Aeromonas aquariorum*		+			+	+
AUN_16	*Aeromonas hydrophila*		+				
AUN_33	*Aeromonas hydrophila*						
AUN_34	*Aeromonas hydrophila*					+	+
AUN_35	*Aeromonas hydrophila*						
AUN_38	*Aeromonas hydrophila*	+	+	+			+
AUN_40	*Aeromonas hydrophila*		+				
CHIN_85	*Aeromonas hydrophila*						+
TAL_129	*Aeromonas hydrophila*	+					+
WAK_2	*Aeromonas hydrophila*						
TAL_55	*Aeromonas jandaei*						
WAK_20	*Aeromonas jandaei*					+	
WAK_91	*Aeromonas jandaei*	+					
TAL_115	*Aeromonas jandaei*	+	+				+
AUN_41	*Aeromonas* sp.	+					+
CHIN_5	*Aeromonas* sp.	+		+			+

Supplementary table 1 (continued)

Isolate code	Identification	*IntI1*	*IntI2*	bla_{SHV}	bla_{OXA}	*aac(6')-Ib-cr*	bla_{CTX-M}
CHIN_72	*Aeromonas* sp.		+				+
TAL_114	*Aeromonas* sp.	+	+				+
WAK_3	*Aeromonas* sp.	+		+			
WAK_4	*Aeromonas* sp.			+			
AUN_31	*Aeromonas sanarellii*					+	+
WAK_93	*Aeromonas sanarellii*	+				+	+
AUN_30	*Aeromonas taiwanensis*						
CHIN_143_a	*Aeromonas veronii*	+		+			
TAL_111	*Aeromonas veronii*	+					+
TAL_113	*Aeromonas veronii*		+			+	
TAL_121	*Aeromonas veronii*					+	+
AUN_14	*Aeromonas veronii*			+			+
TAL_110	*Aeromonas veronii*	+			+	+	+
TAL_50	*Bacillus* sp.			+			
WAK_23	*Bacillus* sp.	+		+			
WAK_136a	*Bacillus firmus*						
WAK_21	*Bacillus flexus*						
AUN_32	*Bacillus flexus*						
WAK_24	*Bacillus flexus*		+				
WAK_94	*Bacillus flexus*						
TAL_51	*Bacillus marisflavi*			+			
TAL_131	*Bacillus megaterium*		+				
AUN_15	*Bacillus megaterium*			+			
CHIN_81	*Bacillus megaterium*					+	+
CHIN_9	*Bacillus megaterium*						
TAL_130	*Bacillus megaterium*	+					
CHIN_88	*Bacillus muralis*	+					+
TAL_106	*Bacillus oceanisediminis*	+		+		+	+
TAL_104	*Bacillus thioparans*		+	+			
CHIN_139	*Enterobacter aerogenes*					+	
WAK_135	*Enterobacter aerogenes*	+				+	
WAK_137	*Enterobacter aerogenes*	+	+			+	
WAK_136	*Enterobacter aerogenes*		+				+
CHIN_84	*Enterobacter asburiae*				+		+
CHIN_86	*Enterobacter asburiae*	+					+
CHIN_147	*Enterobacter asburiae*	+					+
WAK_138	*Enterobacter cloacae*						
AUN_37	*Enterobacter cowanii*			+			+
CHIN_7	*Enterobacter hormaechei*						
CHIN_6	*Enterobacter* sp.						
CHIN_145	*Enterococcus faecium*	+	+		+	+	+
TAL_101	*Escherichia coli*	+				+	+
WAK_96	*Escherichia coli*	+	+	+			
TAL_98	*Shigella dysenteriae*	+			+		
AUN_28	*Shigella dysenteriae*					+	
TAL_99	*Shigella fergusonii*	+	+		+		
AUN_13	*Shigella flexneri*			+		+	

Supplementary table 1 (continued)

Isolate code	Identification	*IntI1*	*IntI2*	bla_{SHV}	bla_{OXA}	*aac(6')-Ib-cr*	bla_{CTX-M}
WAK_22	*Shigella flexneri*			+		+	+
WAK_1	*Klebsiella pneumoniae*						
TAL_102	*Klebsiella pneumoniae*	+		+		+	
CHIN_87	*Leclercia adecarboxylata*	+					
CHIN_8	*Plesiomonas shigelloides*		+	+		+	
CHIN_82	*Plesiomonas shigelloides*			+			
TAL_128_a	*Plesiomonas shigelloides*	+					
TAL_44	*Pseudomonas* sp.	+		+			
TAL_105	*Pseudomonas alcaligenes*		+			+	+
CHIN_89	*Pseudomonas plecoglossicida*			+		+	
TAL_43	*Pseudomonas taiwanensis*			+			
CHIN_74	*Rheinheimera perlucida*						
TAL_45	*Shigella flexneri*			+			
CHIN_142	*Serratia marcescens*					+	+
TAL_120	*Serratia marcescens*	+	+				+
CHIN_143	*Serratia marcescens*			+		+	+
CHIN_144	*Serratia marcescens*	+		+		+	+
CHIN_148	*Serratia marcescens*	+					+
TAL_119	*Serratia marcescens*	+				+	+
TAL_47	*Staphylococcus sciuri*			+			
TAL_52	*Staphylococcus sciuri*				+		
AUN_29	*Vibrio albensis*			+			

+ indicates the presence of the gene, *IntI1*, class 1 integrase, *IntI2*, class 2 integrase, bla_{SHV}, ESBL SHV type; bla_{OXA}, ESBL OXA type; bla_{CTX-M}, ESBL CTXM type; *aac(6')-Ib-cr*, aminoglycoside acetyltransferase; TAL, isolates from Lake Tilapia; AUN, CHIN,WAK, isolates from River Tilapia.