

Supplementary Materials for

Prevalence of Antibiotic Resistance Genes and their Association with Antibiotics in a Wastewater Treatment Plant: Huaguang Liu ¹, Xingyu Zhou ^{2, 3, *}, Hexun Huang ² and Jinsong Zhang ^{1, 2, 3}

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Table S1. Correlation analysis of ARG/16S rRNA gene and corresponding antibiotic residue concentration in collected samples.

	TC	OTC	ΣTCs	ROX	CLA	ΣMLs	NOR	OFX	ΣQNs	SMZ	SMX	ΣSAs	tetA	tetC	tetQ	tetX	tet	sul I	sul II	sul III	sul	intI1
TC	1	0.502	.951**	0.621	0.475	0.559	-0.009	0.501	0.503	-0.193	-0.189	-0.191	-0.053	0.138	-0.422	0.54	0.243	0.626	-0.171	-0.138	0.566	-0.022
OTC	0.502	1	0.744	0.722	0.689	0.718	0.569	0.47	0.483	-0.033	-0.129	-0.101	-0.518	.842*	-0.317	0.595	0.7	-0.106	0.212	0.033	-0.088	0.554
ΣTCs	.951**	0.744	1	0.737	0.612	0.688	0.195	0.554	0.561	-0.161	-0.192	-0.184	-0.226	0.406	-0.439	0.629	0.437	0.446	-0.057	-0.095	0.406	0.18
ROX	0.621	0.722	0.737	1	.934**	.984**	0.405	0.714	0.724	0.102	0.174	0.154	-0.082	0.685	-0.027	0.687	.891**	0.503	0.422	0.319	0.56	0.399
CLA	0.475	0.689	0.612	.934**	1	.982**	0.369	0.624	0.634	-0.077	0.007	-0.018	0.006	0.705	-0.153	.779*	.943**	0.358	0.269	0.274	0.406	0.36
ΣMLs	0.559	0.718	0.688	.984**	.982**	1	0.394	0.681	0.692	0.015	0.095	0.072	-0.04	0.706	-0.09	0.744	.932**	0.44	0.354	0.302	0.493	0.386
NOR	-0.009	0.569	0.195	0.405	0.369	0.394	1	-0.23	-0.212	-0.231	-0.248	-0.245	0.064	.862*	-0.248	-0.178	0.486	-0.113	0.513	0.271	-0.035	1.000**
OFX	0.501	0.47	0.554	0.714	0.624	0.681	-0.23	1	1.000**	0.572	0.59	0.588	-0.486	0.203	0.369	.809*	0.605	0.347	0.263	0.16	0.373	-0.234
ΣQNs	0.503	0.483	0.561	0.724	0.634	0.692	-0.212	1.000**	1	0.57	0.587	0.586	-0.487	0.221	0.366	.809*	0.617	0.347	0.274	0.166	0.374	-0.216
SMZ	-0.193	-0.033	-0.161	0.102	-0.077	0.015	-0.231	0.572	0.57	1	.968**	.984**	-0.579	-0.08	.892**	0.058	0.113	-0.005	0.606	0.19	0.056	-0.218
SMX	-0.189	-0.129	-0.192	0.174	0.007	0.095	-0.248	0.59	0.587	.968**	1	.997**	-0.378	-0.109	.946**	0.066	0.185	0.152	0.627	0.334	0.236	-0.233
ΣSAs	-0.191	-0.101	-0.184	0.154	-0.018	0.072	-0.245	0.588	0.586	.984**	.997**	1	-0.441	-0.101	.936**	0.064	0.164	0.106	0.625	0.293	0.183	-0.23
tetA	-0.053	-0.518	-0.226	-0.082	0.006	-0.04	0.064	-0.486	-0.487	-0.579	-0.378	-0.441	1	-0.213	-0.225	-0.34	-0.108	0.474	-0.175	0.261	0.497	0.073
tetC	0.138	.842*	0.406	0.685	0.705	0.706	.862*	0.203	0.221	-0.08	-0.109	-0.101	-0.213	1	-0.195	0.316	.808*	-0.14	0.488	0.268	-0.064	.855*
tetQ	-0.422	-0.317	-0.439	-0.027	-0.153	-0.09	-0.248	0.369	0.366	.892**	.946**	.936**	-0.225	-0.195	1	-0.138	0.072	0.007	0.539	0.487	0.121	-0.227
tetX	0.54	0.595	0.629	0.687	.779*	0.744	-0.178	.809*	.809*	0.058	0.066	0.064	-0.34	0.316	-0.138	1	0.661	0.187	-0.115	-0.049	0.164	-0.193
tet	0.243	0.7	0.437	.891**	.943**	.932**	0.486	0.605	0.617	0.113	0.185	0.164	-0.108	.808*	0.072	0.661	1	0.165	0.453	0.451	0.262	0.483
sul I	0.626	-0.106	0.446	0.503	0.358	0.44	-0.113	0.347	0.347	-0.005	0.152	0.106	0.474	-0.14	0.007	0.187	0.165	1	0.192	0.062	.980**	-0.109
sul II	-0.171	0.212	-0.057	0.422	0.269	0.354	0.513	0.263	0.274	0.606	0.627	0.625	-0.175	0.488	0.539	-0.115	0.453	0.192	1	0.241	0.274	0.524
sul III	-0.138	0.033	-0.095	0.319	0.274	0.302	0.271	0.16	0.166	0.19	0.334	0.293	0.261	0.268	0.487	-0.049	0.451	0.062	0.241	1	0.255	0.287
sul	0.566	-0.088	0.406	0.56	0.406	0.493	-0.035	0.373	0.374	0.056	0.236	0.183	0.497	-0.064	0.121	0.164	0.262	.980**	0.274	0.255	1	-0.028
intI1	-0.022	0.554	0.18	0.399	0.36	0.386	.988**	-0.234	-0.216	-0.218	-0.233	-0.23	0.073	.855*	-0.227	-0.193	0.483	-0.109	0.524	0.287	-0.028	1

* Significant at the 0.05 level in a two-tailed test. ** Significant at the 0.01 level in a two-tailed test.

