

Supplementary Table S7. Docking results for the phenolic acids, tetracycline, and CCCP against the bacterial *AcrB* efflux pump protein using the SYBYL program.

Compound	Total_ Score	Number of hydrogen bonds	Amino acid residue	Compound	Total_ Score	Number of hydrogen bonds	Amino acid residue
Chlorogenic acid	5.2111	7	GLN-89/GLY-126/SER-128/ARG-620	Gentisic acid	3.0693	4	ASN-68/LYS-163/VAL-175/LEU-177
Sinapic Acid	4.1694	2	GLY-179/ASP-276	α -resorcylic acid	3.0074	6	THR-44/GLN-89/GLU-130/SER-132/GLN-176/LYS-292
Gallic acid	4.0321	5	ASN-68/ASN-70/ASP-174/LEU-177	CCCP	2.9237	1	LYS-770
Vanillic acid	3.9758	3	THR-85/ASP-276/ARG-620	2,3,4-trihydroxybenzoic acid	2.8566	3	GLN-67/ASN-68/LEU-177
Caffeic acid	3.4994	5	ASN-70/LYS-163/VAL-175/LEU-177	2,6-Dihydroxybenzoic acid	2.6898	3	LYS-163/VAL-175/LEU-177
Pyrocatechuic acid	3.497	4	GLN-67/LYS-163/LEU-177	Tetracycline	2.6506	5	ASN-70/LYS-163/GLN-181/LYS-770
<i>M</i> -hydroxybenzoic acid	3.2015	3	ASN-68/ASN-70/LEU-177	Salicylic acid	2.5104	3	LYS-163/VAL-175/LEU-177
β -resorcylic acid	3.1097	5	ASN-68/ASN-70/LYS-163/VAL-175/LEU-177	<i>P</i> -hydroxybenzoic acid	2.4809	3	GLN-151/LEU-177/ALA-279
Protocatechuic acid	3.0761	5	ASN-68/ASN-70/LYS-163/LEU-177				