

Table S1. ECL Tle family phospholipases

Effector class	Bacterial strain	NCBI Refseq	length	predicted ankyrin-repeat immunity protein(s)	
Tle	<i>Enterobacter cloacae</i> subsp. <i>cloacae</i> ATCC 13047	WP_013096193.1	472	WP_013096194.1	
	<i>Enterobacter hormaechei</i> ATCC 49162	WP_006810941.1	488	WP_006810940.1	
	<i>Pseudocherichia vulneris</i> NCTC12130	WP_042387755.1	496	WP_042387753.1; WP_115084734.1	
	<i>Cronobacter sakazakii</i> ATCC BAA-894	WP_007853550.1	476	WP_007889010.1	
	<i>Cronobacter condimenti</i> 1330	WP_007668262.1	488	WP_007668245.1	
	<i>Kosakonia sacchari</i> SP1	WP_017456470.1	472	WP_017456469.1	
	<i>Phytobacter massiliensis</i> JC163	WP_044180714.1	468	WP_237567481.1	
	<i>Lelliottia nimpessuralis</i> 51	TFB20627.1	472	TFB20626.1	
	<i>Citrobacter</i> sp. R56	WP_203360225.1	488	WP_203361220.1	
	<i>Salmonella enterica</i> subsp. <i>enterica</i> serovar <i>Newport</i>	EBR8158188.1	488	EBR8158189.1	
	<i>Franconibacter</i> sp. IITDAS19	WP_024557997.1	479	WP_247365411.1	
	<i>Phytobacter</i> sp. SCO41	WP_108700663.1	474	WP_108700662.1	
	<i>Serratia ureilytica</i> SER00108	WP_197777867.1	493	I5P71_RS11495 (pseudo); WP_197776957.1	
	<i>Serratia marcescens</i> BWH-23	AVU29873.1	493	AVU29874.1; AVU29875.1	
	<i>Serratia marcescens</i> MGH135	WP_033646675.1	493	WP_025302229.1	
	<i>Serratia marcescens</i> Db11	WP_025302228.1	493	WP_025302229.1; WP_025302230.1	
	PAAR	<i>Lonsdalea quercina</i> NCCB 100489	WP_027063756.1	464	WP_026739306.1
		<i>Brenneria</i> sp. L3-3C-1	WP_199379546.1	456	WP_199379544.1; WP_199379560.1; WP_233356511.1
<i>Pectobacterium carotovorum</i> subsp. <i>carotovorum</i> PC1		WP_015841393.1	456	WP_043881824.1	
<i>Dickeya dadantii</i> 3937		WP_013317122.1	456	WP_013317123.1; WP_237703452.1	
<i>Dickeya chrysanthemi</i> Ech1591		WP_012770502.1	456	WP_012770501.1; WP_012770500.1; WP_012770499.1	
<i>Vibrio nigripulchritudo</i> ATCC 27043		WP_004404991.1	513	WP_004404995.1	
<i>Vibrio parahaemolyticus</i> JW16-127		EHA1109881.1	513	EHA1109882.1	
<i>Vibrio mimicus</i> VM223		WP_001138502.1	505	WP_000199146.1; WP_000934380.1	
<i>Polyangium aurulentum</i> SDU3-1		WP_169508841.1	421	WP_136926720.1	
DUF4150 (PAAR-like)	<i>Pantoea</i> sp. PSNIH2	AIX74951.1	422	AIX74950.1	
	<i>Franconibacter pulveris</i> LMG 24059	WP_029591452.1	431	WP_029591453.1	
	<i>Halopseudomonas salegens</i> CECT 8338	WP_092383249.1	434	WP_197675042.1	
	<i>Tahibacter caeni</i> BUT-6	WP_257385295.1	424	WP_257385296.1	
	<i>Xanthomonas translucens</i> pv. <i>cerealis</i>	WP_142742694.1	419	WP_221887383.1	
	<i>Xanthomonas youngii</i> AmX2	WP_206232757.1	433	WP_206232758.1	
	<i>Sorangium</i> sp. Soce836	AUX33661.1	396	AUX33660.1	
	<i>Sorangium cellulosum</i> So0157-2	WP_020737320.1	409	WP_020737319.1	
	<i>Thauera butanivorans</i> NBRC 103042	WP_068639422.1	433	WP_157659236.1; WP_068639420	
	<i>Lampropedia aestuarii</i> YIM MLB12	WP_136407722.1	437	WP_136407723.1; WP_136407723.1	
	<i>Burkholderia vietnamiensis</i> AU35216	WP_249181664.1	382	WP_150331737.1	
	<i>Burkholderia thailandensis</i> Bt10013	WP_150331736.1	400	WP_150331737.1; WP_150331738.1	
	<i>Burkholderia pseudomallei</i> MSHR7500	WP_038799229.1	396	WP_038799227.1	
	VgrG	<i>Aggregatibacter actinomycetemcomitans</i> SL7469	MBN6065491.1	1060	MBN6065490.1
<i>Pasteurella testudinis</i> NCTC12150		WP_115306396.1	1018	WP_115306395.1	
<i>Burkholderia gladioli</i> BSR3		WP_013700226.1	996	WP_013700225.1	
<i>Burkholderia multivorans</i> AU16734		WP_105792066.1	1001	WP_059936093.1	
<i>Burkholderia cepacia</i> ATCC 25416		WP_027786768.1	999	WP_226153282.1	
<i>Desulfuromonas versatilis</i> NIT-T3		WP_221250508.1	1082	WP_221250509.1	
<i>Chondromyces crocatus</i> Cm c5		WP_063796395.1	1019	WP_050434209.1	
<i>Geobacter benzoatilyticus</i> Jerry-YX		WP_207164315.1	942	WP_207164316.1	
<i>Cystobacter fuscus</i> DSM 52655		WP_157758952.1	1050	WP_157758953.1	
Rhs		<i>Pseudomonas fluorescens</i> UM270	WP_042727971.1	4024	WP_080921999.1
	<i>Pseudomonas palleroniana</i> PS006	WP_060752379.1	3668	WP_158510675.1	
	<i>Vibrio vulnificus</i> NBRC 15645 = ATCC 27562	WP_017422738.1	1515	WP_017422739.1	
	<i>Vibrio cincinnatiensis</i> 19-VB00020	WP_238132309.1	1558	WP_238124837.1	
CdiA	<i>Desulfovibrio subterraneus</i> ND17	WP_269890085.1	1965	WP_269890084.1	
	<i>Cupidesulfovibrio liaohensis</i> XJ01	WP_167127856.1	1900	WP_167127853.1 (partial)	
	<i>Desulfomicrobium</i> sp. ZS1	WP_254080456.1	2300	WP_254080455.1	
	<i>Pseudodesulfovibrio piezophilus</i> C1TLV30	CCH47381.1	1774	CCH47382.1	