Supplemental Table S1 Predicted *Burkholderia*-type CDI system-encoding loci

Locus	BcpA[1]	LYN[2]	BcpI[1]	BcpO[1]	BcpB Acc. No.[3]	Identical Alleles
B. thailandensis E264†	3147	lyn	270	74	YP_443235.1	N/A
B. thailandensis TXDOH†	3025	lyn	171	73	ZP_02369419.1	N/A
B. pseudomallei 1106A-1†	3123	lyn	99	73	YP_001068100.1	*Bp*BCC215-2, *Bg*BSR3-1
B. pseudomallei 1106A-2†	3131	lyn	117	74	YP_001065412.1	*Bp*B7210-2, *Bp*1710b-1, *Bp*112-2, *Bp*Pasteur52237
B. pseudomallei 1106A-3†	3141	lyn	109	57*	YP_001076829.1	*Bp*DM98, *Bp*S13, *Bp*1710b-2, *Bp*B7210-1, *Bp*1655, *Bp*112-1
B. pseudomallei 1710b-1†	3131	lyn	117	74	YP_332689.1	*Bp*1106A-2, *Bp*B7210-2, *Bp*112-2, *Bp*Pasteur52237
B. pseudomallei 1710b-2†	3141	lyn	109	57*	YP_336326.1	*Bp*DM98, *Bp*S13, *Bp*1106A-3, *Bp*B7210-1, *Bp*1655, *Bp*112-1
B. pseudomallei 668†	3134	lyn	118	74	YP_001058175.2	N/A
B. pseudomallei K96243†	3103	lyn	109	172	YP_112054.1	*Bp*91
B. pseudomallei 1026b	3122	lyn	101	81	–	N/A
B. pseudomallei 112-1	3141	lyn	109	57*	ZP_02503380.1	*Bp*DM98, *Bp*S13, *Bp*1710b-2, *Bp*1106A-3, *Bp*B7210-1, *Bp*1655
B. pseudomallei 112-2	2839	lyn	117	74	ZP_02497140.1	*Bp*1106A-2, *Bp*B7210-2, *Bp*1710b-1, *Bp*Pasteur52237
B. pseudomallei 14	3057	lyn	166	–	ZP_02416760.1	N/A
B. pseudomallei 1655	3141	lyn	109	57*	ZP_04891151.1	*Bp*DM98, *Bp*S13, *Bp*1710b-2, *Bp*1106A-3, *Bp*B7210-1, *Bp*112-1
B. pseudomallei 305	3107	lyn	96	74	ZP_01767849.1	*Bp*MSHR346, *Bp*7894-1, *Bp*406e, *Bp*576-2
B. pseudomallei 406e	3107	lyn	96	74	ZP_04965785.1	*Bp*MSHR346, *Bp*305, *Bp*7894-1, *Bp*576-2
B. pseudomallei 576-1	3083	lyn	99	81	ZP_03453873.1	N/A
B. pseudomallei 576-2	3107	lyn	96	74	ZP_03451717.1	*Bp*MSHR346, *Bp*305, *Bp*7894-1, *Bp*406e
B. pseudomallei 7894-1	2932	lyn	96	74	ZP_02487784.1	*Bp*MSHR346, *Bp*305, *Bp*406e, *Bp*576-2
B. pseudomallei 7894-2	3125	lyn	88	73	ZP_02483696.1	N/A
B. pseudomallei 7894-3	2980	lyn	83	–	ZP_02487474.1	N/A
B. pseudomallei 91	3103	lyn	109	172	ZP_02452844.1	*Bp*K96243
B. pseudomallei B7210-1	3141	lyn	109	57*		*Bp*DM98, *Bp*S13, *Bp*1710b-2, *Bp*1106A-3, *Bp*1655, *Bp*112-1
B. pseudomallei B7210-2	2956	lyn	117	74	ZP_02470328.1	*Bp*1106A-2, *Bp*1710b-1, *Bp*112-2, *Bp*Pasteur52237
B. pseudomallei B7210-3	3223	lyn	97	–	ZP_02477263.1	N/A
B. pseudomallei BCC215-1	3122	lyn	101	147	ZP_02511212.1	N/A
B. pseudomallei BCC215-2	3123	lyn	99	73	ZP_02511816.1	*Bp*1106A-1, *Bg*BSR3-1
B. pseudomallei DM98	3141	lyn	109	57*	ZP_02408285.1	*Bp*1106A-3, *Bp*S13, *Bp*1710b-2, *Bp*B7210-1, *Bp*1655, *Bp*112-1
B. pseudomallei MSHR346	3107	lyn	96	74	YP_002895846.1	*Bp*305, *Bp*7894-1, *Bp*406e, *Bp*576-2
B. pseudomallei NCTC13177	3096	lyn	111	81	ZP_02495130.1	N/A
B. pseudomallei Pasteur52237	3131	lyn	117	74	ZP_04897242.1	*Bp*1106A-2, *Bp*B7210-2, *Bp*1710b-1, *Bp*112-2
B. pseudomallei S13	2850	lyn	109	57*	NZ_CH899763.1	*Bp*DM98, *Bp*1710b-2, *Bp*1106A-3, *Bp*B7210-1, *Bp*1655, *Bp*112-1

B. ambifaria AMMD[†]	3028	lyn	155	89	YP_775670.1	N/A
B. ambifaria IOP40-10	3066	lyn	89**	–	ZP_02889733.1	N/A
B. ambifaria MEX-5	3776	lyn	128	99	ZP_02907827.1	N/A
B. cenocepacia H111	3229	–	142	59	CCE53449.1	N/A
B. dolosa AUO158-1	3379	lyn	270	89**	ZP_04945975.1	N/A
B. dolosa AUO158-2	3037	lyn	164	67**	ZP_04947043.1	N/A
B. gladioli BSR3-1	3123	lyn	99	73	YP_004358894.1	*Bp*1106A-1, *Bp*BCC215-2
B. gladioli BSR3-2(p)	3082	lyn	103	–	YP_004350998.1	N/A
B. gladioli BSR3-3	3038	lyn	77	113	YP_004349107.1	N/A
B. glumea BGR1[†]	3151	lyn	269	71	YP_002909358.1	N/A
B. multivorans CGD2M-1	3060	lyn	93	118	ZP_03577349.1	N/A
B. multivorans CGD2M-2	3423	venn	106	–	ZP_03575068.1	N/A
B. phymatum STM815-1[†]	3079	–	44	–	YP_001860455.1	N/A
B. phymatum STM815-2(p)[†]	3015	–	187	54	YP_001863347.1	N/A
B. rhizoxinica HKI454[†]	2981	–	83	–	YP_004028152.1	N/A
B. ubonensis BU	2814	lyn	89	56	ZP_02379486.1	N/A
B. vietnamiensis G4[†]	3513	lyn	81	64	YP_001116443.1	N/A
B. xenovorans LB400[†]	2984	–	81	83	YP_557396.1	N/A
R. solanacearum CFBP2957(p)[†]	3651	–	301	143	YP_003747808.1	N/A
R. solanacearum CMR15(p)[†]	3552	–	145	59	CBJ40172	N/A
R. solanacearum GMI1000[†]	3501	lyn	80	155	NP_519009.1	N/A
R. solanacearum Po82-1[†]	3305	venn	182*	179	AEG67591.1	N/A
R. solanacearum Po82-2(p)[†]	3177	–	190	76	AEG71437.1	N/A
R. solanacearum PSI07[†]	3386	–	166	51	YP_003750973.1	N/A
R. syzygii R24	3208	–	186	86	CCA87914.1	N/A
C. metallidurans CH34(p)[†]	3004	–	77	63	YP_587468.1	N/A

[1]Amino acid length of predicted protein

[2]Nx(E/Q)LYN (or VENN) motif in predicted BcpA sequence located ~350 amino acid from the C-terminus of the protein

[3]Accession number in NCBI database

[†]Loci from complete genome sequences

*Encoding ORF overlaps the 3' ORF by ~50-100 nucleotides

**Encoding ORF begins with a 'gtg' sequence

(p) Locus located on plasmid

N/A Allele not found in any other strain