

S5 Table. Genetic diversity, geographic distribution and clinical outcomes of melioidosis patients with *Burkholderia pseudomallei* that shared the founder genotype ST1132.

	Isolate		YLF/ BTFC clade	<i>fhab3</i>	<i>bimA_{BM}</i> / <i>bimA_{BP}</i>	LPSA	Geographical region	Clinical presentations
1	BPs2a/b		YLF	+	<i>bimA_{BP}</i>	+	CP	multiple skin and soft tissue abscesses, septic arthritis, sepsis
2	BPs11		YLF	+	<i>bimA_{BP}</i>	+	WP	psoas abscess
3	BPs43		YLF	+	<i>bimA_{BP}</i>	+	UVA	multiple skin and soft tissue abscesses, septic arthritis, sepsis
4	BPs45		YLF	+	<i>bimA_{BP}</i>	+	NWP	abscess over the chest, P/H of abscess over right thigh
5	BPs60		YLF	+	<i>bimA_{BP}</i>	+	NWP	sepsis/ headache / abdominal pain
6	BPs61		YLF	+	<i>bimA_{BP}</i>	+	NWP	splenic abscess, sepsis
7	BPs64		YLF	+	<i>bimA_{BP}</i>	+	NWP	R/S knee joint septic arthritis
8	BPs91		YLF	+	<i>bimA_{BP}</i>	+	NWP	endocarditis of aortic valve
9	BPs 144		YLF	+	<i>bimA_{BP}</i>	+	NWP	sepsis
10	BPs 155		YLF	+	<i>bimA_{BP}</i>	+	WP	sepsis
11	BPs 171		YLF	+	<i>bimA_{BP}</i>	+	WP	NA
12	BPs 180		YLF	-	<i>bimA_{BP}</i>	+	NWP	cervical lymph node abscess
13	BPs 186		YLF	+	<i>bimA_{BP}</i>	+	WP	sepsis, prostatitis,
14	BPs 191		YLF	-	<i>bimA_{BP}</i>	+	NWP	parotid abscess
15	BPs 196		YLF	-	<i>bimA_{BP}</i>	+	NWP	splenic abscess
16	BPs 204		YLF	+	<i>bimA_{BP}</i>	+	NWP	parotid abscess

UVA - Uva Province, NWP - North Western Province, CP - Central Province, WP - Western Province, SP - Southern Province, NCP - North Central Province, SGP - Sabaragamuwa Province, *bimA_{BP}*/*bimA_{BM}* – *Burkholderia* intracellular motility factor (*BimA*) gene variants, *fhaB3* - filamentous hemagglutinin 3, YLF – *Yersinia*-like fimbrial gene cluster (YLF-clade), BTFC - *Burkholderia thailandensis* flagellum and chemotaxis gene cluster (BTFC-clade), LPSA - lipopolysaccharide (LPS) O-antigen type A, + - positive, - - negative, nd - not determined.