

Isolate	<i>fliC</i> Probe G-T Mismatch 1 FoD: 0.997	<i>fliC</i> Probe C-C Mismatch 2 FoD: 0.985	<i>fliC</i> Forward C-C Mismatch FoD: 0.940
<i>Mismatch profile 1</i>			
Bp TSV202	x		
Bp MSHR 1655	x		
<i>Mismatch profile 2</i>			
Bt 34		x	
Bt 2003015869		x	
Bt H0587		x	
<i>Mismatch profile 3</i>			
Bo C6786		x	x
Bo EO147		x	x

FoD, Fraction of duplex; Bp, *B. pseudomallei*; Bt, *B. thailandensis*; Bo, *B. oklahomensis*