

<b>Isolate</b>	<b><i>orf11</i> Forward T-G Mismatch 1 FoD: 0.991</b>	<b><i>orf11</i> Forward G-G Mismatch 2 FoD: 0.926</b>	<b><i>orf11</i> Reverse A-C Mismatch FoD: 0.989</b>	<b><i>orf11</i> Probe G-T Mismatch FoD: 0.992</b>
<i>Mismatch profile 1</i>				
Bp HB PUB10134a	x		x	
Bp PHL 112*	x		x	
<i>Mismatch profile 2</i>				
Bp BDP		x	x	
Bp MSHR 305		x	x	
Bp MSHR 520		x	x	
<i>Mismatch profile 3</i>				
Bp Bp1651			x	
Bp K42			x	
Bp MSHR146*			x	
Bp MSHR 346			x	
Bp MSHR 511			x	
Bp PB08298010			x	
Bp NAU20B-16			x	
Bp TSV48			x	
<i>Mismatch profile 4</i>				
Bp MSHR 391			x	x
Bp MSHR 491			x	x
Bp MSHR 668			x	x

FoD, Fraction of duplex; Bp, *B. pseudomallei*; Bt, *B. thailandensis*; \*, indicates sample was previously tested by the *Bcom* assay