

**Table S2. Primers used in this study**

Cloning <i>hrpA</i> in pJET1.2			
B1601-NdeI-catATGAATGATTTCAAACCTC <sup>1</sup>			
B1603-BamHI-ggatccATAACAAGGCTTCAAAGTTAG <sup>1</sup>			
Site Directed Mutations in <i>hrpA</i>			
Mutation Point	Primers		
D126A	B1886-GAATATGATGTAATAATAATAGCCGAAGCACACGAAAGAAG B1895-CTTCTTTTCGTGTGCTTCGGCTATTATTATTACATCATATTC		
E127A	B1888-GATGTAATAATAATAGACGCAGCACACGAAAGAAGTTAAAC B1889-GTTTAAACTTCTTTCGTGTGCTGCGTCTATTATTATTACATC		
S158A	B2185-GATTTTAAAATCATAGTTTCGGCTGCTACAATAAACACAAAAA B2186-TTTTTGTGTTTATTGTAGCAGCCGAAACTATGATTTTAAATC		
T160A	B2187-CATAGTTTCGTCTGCTGCAATAAACACAAAAATATT B2188-AATATTTTTGTGTTTATTGCAGCAGACGAAACTATG		
I285A	B2183-CAACATAGCAGAAACTTCAGCCACAATTGAAAATATTAAAT B2184-ATTTTAATTTTCAATTGTGGCTGAAGTTTCTGCTATGTTG		
Complementation plasmid construction			
Annealing site	Primers <sup>2</sup>		
<i>hrpA</i>	B1950-BamHI-cgcgatccgATGAATGATTTCAAACCTCCAAT <sup>1</sup>		
<i>PflgB-kan</i> + <i>hrpA</i>	B1951-tctcctgaagctcggtatTAAAAAAGTAATTAATTTTTGTAA		
<i>hrpA</i> + <i>PflgB-kan</i>	B1952-TTACAAAAAATTAATTTACTTTTTTAAatcccgagctcaaggaaga		
between <i>hrpA</i> and <i>bb0826</i> + <i>PflgB-kan</i>	B1953-TCAAAGTTTACTTTTTTAAAAAAGTAttagaaaaactcatcgagcatc		
<i>PflgB-kan</i> + between <i>hrpA</i> and <i>bb0826</i>	B1954-gatgctgatgagttttctaaTACTTTTTTAAAAACTAAACTTTGA		
<i>bb0826</i>	B1955-EcoRI-ccggaattccggtTATTCTTTCTTTTTTATAAGA <sup>1</sup>		
Screening <i>hrpA</i> mutants			
	Annealing site	Primers <sup>3</sup>	
Point mutations in <i>hrpA</i>	D126A	B2176-CGTGTGCTTCGG	
	E127A	B2173-TTCTTTCGTGTGCTG	
	S158A	B2201-GTTTATTGTAGCAGC	
<i>hrpA</i> complementation	<i>hrpA</i>	B1219-GTTATTTTTGTATTCGGCTTT <sup>4</sup> B1220-TTCGGCTGCTACAATAAACAC <sup>4</sup>	
	$\Delta$ <i>hrpA</i>	B1398-TTAAAACTTCAAAGATATTAACAA B1399-GCAGGAAGACTTTCAAAA	
	<i>PflgB-aacC1</i> (gent)	B348-CGCAGCAGCAACGATGTTAC B349-CTTGACGTAGATCACATAAGC	
	<i>PflgB-kan</i>	B70-CATATGACCATATTCAACGGGAAACG B71-AAAGCCGTTTCTGTAATGAAGGAG	
Northern Blotting Probes			
Target Transcript	Primers		
<i>bb0241</i>	B2205-GAATTATGCTTTGGAACAATAG B2206-GGCACAAAAATAAACCACC		
	B2221-GATTTTTTGAACTAAAAACAAC B2222-GAATAGCACTATAGACCAATTG		
<i>bb0242</i>	B2223-GCTCTGTTCTATATTACGATGATTC B2224-GTTTCTTGAGATTTTTGAAGTG		
	B2225-CAAGAGAATGACAAAGACACTC B2226-CCAAAGAGTCTTGTACTGTAGG		
<i>bb0243</i>	B2227-GTTTGCAGATTCTAACAATGC B2228-CTTTGCAACATCTTTTTTTG		
	B2227-GTTTGCAGATTCTAACAATGC B2228-CTTTGCAACATCTTTTTTTG		
<i>bb0603</i>	B2225-CAAGAGAATGACAAAGACACTC B2226-CCAAAGAGTCTTGTACTGTAGG		
	B2227-GTTTGCAGATTCTAACAATGC B2228-CTTTGCAACATCTTTTTTTG		
<i>bb074</i>	B2227-GTTTGCAGATTCTAACAATGC B2228-CTTTGCAACATCTTTTTTTG		
	B2227-GTTTGCAGATTCTAACAATGC B2228-CTTTGCAACATCTTTTTTTG		

<sup>1</sup> Lower case letters indicate sequences added for the indicated restriction site.

<sup>2</sup> Upper and lower case letters indicate sequence from *B. burgdorferi* genome and *PflgB-kan* resistance cassette, respectively. Except where indicated <sup>1</sup>.

<sup>3</sup> Bold characters are the mutation points in the wild-type *hrpA* sequence.

<sup>4</sup> Primer used in Salman-Dilgimen *et al.* 2011