

Table S1. List of primers used in the multilocus sequencing typing (MLST) scheme according to Margos et al. [29] for amplification of eight genes of *Borrelia* spp. of the relapsing fever group.

Gene	Primers	DNA sequences (5'-3') of forward (F) and reverse (R) primers of each pair of primers	Amplicon size (bp)	Reference
<i>clpA</i>	clpAF1262	F- AAGCTTTGAYYTATTAGATGGTC	1014	http://pubmlst.org/borrelia
	clpAR2276	R- TCATATTTDATRGTDTCGTC		
	clpAF1264	F- GCTTTGAYYTATTAGATGGTC	834*	
	clpAR2098	R- CAAAAAAAASATCHARRTTTCATTTTTAG		
<i>rplB</i>	rplF3	F- GGAGAAAAATATGGKATTAAGAC	766	http://pubmlst.org/borrelia
	rplR769	R- GRCCCCAAGGWGATAC		
	rplF18	F- ATTAAGACTTATARGCCAAAAAC	741*	
	rplF759	R- GATACAGGATGWCAGCC		
	rplF24	F- GACTTATARGCCAAAAACTTC	741*	
	rplF765	R- CCAAGGWGATACAGGATG		
<i>pyrG</i>	pyrF372	F- TGGRAARTATTAGGKAGAACTG	1003	http://pubmlst.org/borrelia
	pyrR1375	R- TATTAGGKAGAACTGTACAGC		
	PyrF379	F- TATTAGGKAGAACTGTACAGC	873*	
	PyrR1252	R- TATRCCACAAACATTACGKGC		
<i>recG</i>	recF898	F- GCKTTCTMTCTAGYATTCC	881	http://pubmlst.org/borrelia
	recR1779	R- TCRGTTAAAGGTTCTTATAAAG		
	recF908	F- CTAGYATTCTYTAATTGARGC	871*	
	recR1779	R- TCRGTTAAAGGTTCTTATAAAG		
<i>clpX</i>	clpXF109	F- GCYATTGTTTGAAATGTTCTAAATATG	859	http://pubmlst.org/borrelia
	clpXR1277	R- TAAAGTTCTTGCCCAAGG		
	clpXF268	F- GCYATTGTTTGAAATGTTCTAAATATG	721*	
	clpXR1183	R- CTTTTTAATTGCTTASTWGAAGG		
<i>pepX</i>	pepXF361	F- AGAGAYTTAAGYTTAKCAGG	1168	http://pubmlst.org/borrelia
	pepXR1207	R- CYATAGTTCTTAAAGAYTGC		
	pepXF364	F- GAYTTAAGYTTAKCAGGARTTG	823*	
	pepXR1187	R- TGCATTCCCCACATTGG		
<i>nifS</i>	nifF23	F- TAAAAATAAAGAGTACTAAATAATG	876	http://pubmlst.org/borrelia
	nifR899	R- CCAAGACCAATAATTCTGC		
	nifF60	F- AAATAATGGTATAATTCTAAATAA	656*	
	nifR716	R- GGAGCAAGCATTTYTGTC		
<i>uvrA</i>	uvrF1170	F- GAGGCATTATCTTWCAAC	983	http://pubmlst.org/borrelia
	uvrR2153	R- CTAATCTCDGTAAAAATCCAACATAAG		
	uvrF1173	F- GCGTTATCTTWCAACTGAATC	974*	
	uvrR2147	R- TCDGTAAAAATCCAACATAAGTTGC		

* amplified through a nested or heminested reaction.