

Table S1. Primers for cloning LigB proteins.

<i>Forward primers</i> ^a	
7'	TATAAAC <u>CATATGG</u> GCTGAAATTA AAAATACCAGTGGAAG
8	TATTTAC <u>CATATG</u> CAATTGATTTCCATTGCCGTA
P-8 ^b	CAATTGATTTCCATTGCCGTA
9	ATACTCC <u>CATATG</u> CTTCTTACTTCTATCGAGATAACACCG
9' ^c	CTGCTT <u>CATATG</u> TCTTCCGATCCATCTAAGATTG
10	ACCGTAC <u>CATATG</u> AAACTGAAAAGTATAACTATCAGTCCTTCC
11	ATAC <u>CATATG</u> ACGTTAGATTCCATTA AAATCAATCCAGT
11' _{Pom} ^d	GATCGAC <u>CATATG</u> ACTGCGACTTACAATTCCATC
P-12 ^b	ACCCTTTCTTCGATTTCAATATCTCCTATCAA
<i>Reverse primers</i> ^e	
8	TAAAAT <u>CTCGAG</u> TGCAGGAGTGACATTCAAAC
9	TTTAAT <u>CTCGAG</u> TAAGTCAGTGACTGTAATCGGAATTG
10	ATTTTACT <u>CTCGAG</u> GGCAGCACTTACATTA AAATTTATTTTA
P-10 ^b	GGCAGCACTTACATTA AAATTTATTTTATTACTGCTTATAG
11	ATATTT <u>CTCGAG</u> TGCTGCGCTGACCGTTATG
11' ^f	CTGTTACT <u>CTCGAG</u> ACTGGACCAGGTA ACTGAATC
12	ATTTAACT <u>CTCGAG</u> AGCTATCGTGTCCGTTTTGTTTAC
CTD _{Pom} ^g	CTATTT <u>CTCGAG</u> TTAATTGGA ACTATTAATTATTTTGTAATTGG

^aForward primers are named for the respective LigB repeats as in **Table 1** and displayed 5' to 3' with leaders containing an *Nde* I restriction endonuclease site that is underlined. See **Table 1** for the initial amino acid of each LigB repeat from *L. interrogans* serovar Copenhageni strain Fiocruz L1-130.

^b5'-phosphorylated for blunt-end ligation to 3'-end of linker-encoding sequence following repeat 10.

^cN-terminal-deleted repeat 9 starting at S803.

^dN-terminal-deleted repeat 11 from *L. interrogans* serovar Pomona starting at T1014.

^eReverse primers contain an *Xho* I site that is underlined. See **Table 1** for the last residue of the linker sequence that follows each repeat except for repeat 12.

^fC-terminal-deleted repeat 11 ending at S985.

^gFirst 45 amino acids of the C-terminal domain in the Pomona strain ending at N1165.