

Table S2. Oligonucleotides used in this study

Oligonucleotides	Sequence ^a (5'-3')	Localization ^b	Restriction site
<i>pilC1_EcoRI_F</i>	CCG <u>gaattc</u> CAGTAAATACGCTATTATC	1952796 - 1952814	<i>EcoRI</i>
<i>pilC1_XbaI_R</i>	CCG <u>tctaga</u> TTTCAATATCCGACTTAT	1953907 - 1953890	<i>XbaI</i>
<i>pilC1_invF</i>	CGG <u>cctagg</u> CGCCAAACTGCACCTGC	1953462 - 1953478	<i>AvrII</i>
<i>pilC1_invR</i>	CGG <u>cctagg</u> CCCGCATTTTTTACCAAG	1953206 - 1953189	<i>AvrII</i>
<i>pilC2_EcoRI_F</i>	CGG <u>gaattc</u> CCTATCCATACGTTATTG	44683 - 44700	<i>EcoRI</i>
<i>pilC2_XbaI_R</i>	CGG <u>tctaga</u> TTTTAATCTGCGTATCAC	45831 - 45814	<i>XbaI</i>
<i>pilC2_invF</i>	CGG <u>cctagg</u> GTCTTTTATCTGAACGCC	45372 - 45389	<i>AvrII</i>
<i>pilC2_invR</i>	CGG <u>cctagg</u> TATATATCCAGCTTGCCG	45118 - 45101	<i>AvrII</i>
<i>pilE_Xho_F</i>	GCGCGC <u>cctcgag</u> ATGAACACCCTTCAAAAAGG	17495 - 17476	<i>XhoI</i>
<i>pilE_Xba_R</i>	GCGCGC <u>tctaga</u> TTAGCTGGCAGATGAATC	16989 - 17006	<i>XbaI</i>
<i>pilE_inv_F</i>	CTGTGG <u>cctagg</u> CGTCAAAACGGTTCGGTC	17144 - 17127	<i>AvrII</i>
<i>pilE_inv_R</i>	ACGGCT <u>cctagg</u> TGACCTTCAGCCAAAAGG	17347 - 17364	<i>AvrII</i>
<i>pilC1_exp_F</i>	ACTTGTCTATAAAACCCG	1953150 - 1953167	
<i>pilC1_exp_in_R</i>	CCCAAGGAAATATCGCGG	1953419 - 1953402	
<i>pilC1_exp_out_R</i>	GGTGATATCTTTAATCCC	1953507 - 1953490	
<i>pilC2_exp_F</i>	AACCCAATTTACCTTCGG	45056 - 45073	
<i>pilC2_exp_in_R</i>	GTATGGGCTTTGCGCACC	45290 - 45273	
<i>pilC2_exp_out_R</i>	ACTGTTATGCCTTGCGCG	45445 - 45428	
<i>pilE_exp_F</i>	GGTTTTACCCTTATCGAGCT	17477 - 17458	
<i>pilE_exp_R</i>	GGATGGCTTCGGAAACTTG	17363 - 17381	

^arestriction sites are underlined, ^bcoordinates according to genome annotation from (4).