

Oligonucleotides	Sequences	Amplicons and comments
bvgS-Up bvgS-Lo	5'- TTCTAGACATTGCGAAGATCG -3' 5'- ATAAGCTTCATGGTCAGCCTCAGGTCC -3'	526 bp <i>Xba</i> I- <i>Hind</i> III <i>bvgS</i> fragment for construction of BPSM <sub>ΔbvgAS</sub> and allelic exchange
bvgA-Up bvgA"-Lo	5'-TAGAATTCAACGGCATTGACGGGCTCA-3' 5'-TAGGTACCCTAGTTGGACAACACGCTGATGA-3'	504 bp <i>Eco</i> RI- <i>Kpn</i> I <i>bvgA</i> fragment for construction of BPSM <sub>ΔbvgAS</sub>
bvgA-Lo	5'-ATACTAGTGACGTGGCGTTCAGCTTCTG-3'	504 bp <i>Eco</i> RI- <i>Spe</i> I <i>bvgA</i> fragment for allelic exchange (used with bvgA-Up)
bvgA'-Up bvgA'-Lo	5'-AGCCATTCTTGTACGCATCA-3' 5'-CAGATCAGACTGCGGGGGTAC-3'	762 bp fragment coding for BvgA (sequencing)
VFT1-Up VFT1-Lo	5'-ATACTAGTGGAACTGATAGACCTCGCAA-3' 5'-TAGGATCCCAGTCCCCAGCGGTAGATCA-3'	904 bp <i>Spe</i> I- <i>Bam</i> HI fragment coding for VFT1 (cassette exchange)
VFT1'-Up VFT1'-Lo	5'-GCAGAACGCTGAACGCCACGTC-3' 5'-ATTCAGGACCGCCACCTTGAC-3'	1020 bp fragment coding for VFT1 (sequencing)
VFT2-Up VFT2-Lo	5'-TAGGATCCAGCATTGCGCTCGATTG-3' 5'-GGTAGATCTCGTTGCGGTAGG-3'	786 bp <i>Bam</i> HI- <i>Bgl</i> II fragment coding for VFT2 (cassette exchange)
VFT2'-Up VFT2'-Lo	5'-GCCTGCTGCGGGTGGTCA-3' 5'-TCGATGAGCACCGCGCATGAAT-3'	1000 bp fragment coding for VFT2 (sequencing)
PAS/HisKin-Up ΔEcoRI-Lo ΔEcoRI-Up PAS/HisKin-Lo	5'-ACCGAACGAGATCTACCTG-3' 5'-AAATTCCAGCTGGTCGTT-3' 5'-GAACGACCAGCTGGAATTATGCGCGTGCTCATCGACGG-3' 5'-TATCTAGAATGTCGCCTATCAGC-3'	708 bp <i>Bgl</i> II- <i>Xba</i> I fragment coding for the PAS and Histidine Kinase domains (overlapping PCR to remove EcoRI site from the cassette). PAS/HisKin-Up and Lo also used for sequencing
R-Up R-Lo	5'-CCGGGCCAAGACCACGTTCC-3' 5'-TGCACCGCGTACGTCCGTAAAC-3'	1007 bp fragment coding for receiver domain of BvgS (sequencing)
Hpt-Up Hpt-Lo	5'-ACTACCTGGGCCAGCGTGTC-3' 5'-CGGTATCTGCTGGCGACCAT-3'	879 bp fragment coding for Hpt domain of BvgS (sequencing)