

Table 1S
Oligonucleotides used

SrtAF	5' end of <i>srtA</i> of <i>B. anthracis</i> gacagAAGCTT <u>aaggagtgaaggtttgtat</u> gaataagcaaagaatttata
SrtAR	3' end of <i>srtA</i> of <i>B. anthracis</i> gacagGGATCC <u>ttatattatcatcatcatctttataat</u> cgttcgc
PhrCF	5' end of <i>phrC</i> of <i>B. subtilis</i> gacagAAGCTT <u>aaggagtgaaggtttgtat</u> g
PhrCR	3' end of <i>phrC</i> of <i>B. subtilis</i> gacagACTAGTgtcacatgaaagtgcgtg
HisCelAF	5' end of <i>celA</i> of <i>C. thermocellum</i> gacagACTAGT <u>catcatcatcatcatcat</u> gcagggtgcctttacacaaaatac
HisCelAR	3' end of <i>celA</i> of <i>C. thermocellum</i> gacagGCGGCCGCa <u>taaggtaggtgggtatgc</u>
FibF	5' end of the c-terminal domain of <i>fibronectin binding protein B</i> of <i>S. aureus</i> gacagGC <u>GGCCG</u> Ccaaga <u>tgttgtataatgaa</u> ggtc
FibR	3' end of the c-terminal domain of <i>fibronectin binding protein B</i> of <i>S. aureus</i> gacagGCATG <u>Cttatgc</u> ttgtgattctttat <u>ttctgc</u>
CohF	5' end of the second cohesin of <i>cipA</i> of <i>C. thermocellum</i> gacagACTAGT <u>catcatcatcatcat</u> gggtggtagaaattgg
CohR	3' end of the second cohesin of <i>cipA</i> of <i>C. thermocellum</i> gacagGC <u>GGCCG</u> C <u>tttgtcggtgtgc</u> atgc
DoctF	5' end of the CBM and dockerin module of <i>xyn10B</i> of <i>C. thermocellum</i> gacagGC <u>GGCCG</u> C <u>CC</u> c <u>agtcc</u> aatggcgacggtaa
DoctR	3' end of the CBM and dockerin module of <i>xyn10B</i> of <i>C. thermocellum</i> gacagGCATG <u>Ctaaggat</u> ttctgctacagg
CelA-DoctF	5' end of the <i>celA-doct</i> fusion protein gacaGCTAGCgcagggtgtgc <u>cttt</u> aacac
CelA-DoctR	3' end of the <i>celA-doct</i> fusion protein gacagAAGCTT <u>taaggat</u> ttctgctacag
GSTF	5' end of <i>gst</i> of pGEX-4t containing 20 nt from <i>fibronectin binding protein B</i> of <i>S. aureus</i> gaaataaaaagaat <u>acaaagcaatgtcccataactaggttat</u> gg
GSTR	3' end of <i>gst</i> of pGEX-4t gacagGCATG <u>Cttagt</u> cacatgc <u>ggccg</u> c <u>ctcg</u>
FibGSTR	3' end of <i>fibronecting binding protein B</i> of <i>S. aureus</i> containing 20 nt of <i>gst</i> of pGEX-4t ccaataac <u>cttagtatagggacat</u> tgcttgcattttttat
Scaff	5' end of fusion scaffold containing a type I cohesin of <i>C. cellulolyticum cipC</i> , a type I cohesin and CBM of <i>C. thermocellum cipA</i> , and a type I cohesin of <i>scaB</i> of <i>R. flavefaciens</i> gacagACTAGT <u>taccat</u> ac <u>gtgtcc</u> agattac <u>gt</u> ggcgatt <u>ctt</u> aa <u>gttac</u> ag
ScafR	3' end of fusion scaffold containing a type I cohesin of <i>C. cellulolyticum cipC</i> , a type I cohesin and CBM of <i>C. thermocellum cipA</i> , and a type I cohesin of <i>scaB</i> of <i>R. flavefaciens</i> gacagGC <u>GGCCG</u> C <u>tttacaa</u> at <u>gtatgc</u> cc

Restriction sites are in uppercase. Ribosomal binding sites used are in bold. Nucleotide sequence for FLAG tag is italicized and underlined. Nucleotide sequence for polyhistidine tag is underlined. Nucleotide sequence for HA tag is italicized.