

Supplementary table 1. Oligonucleotides used in this study

Oligonucleotide	Sequence (5'-3') ^a	Use in this work
Generation of CipA variants with single aa substitutions, deletions in various regions, and extensions		
CipA-359(-)	TTTTTAGGCTTTGTCGACTTCTTAT ACAACCATATCTAACACATC	Generation of CipA ₁₉₋₃₅₉ (aa 19-359), C-terminal deletion of aa 359-369
CipA-145(+)	GCATATTCCAAGGATCCAAAGCAA CTAATACTCAAGC	Generation of CipA ₁₄₅₋₃₆₉ (aa 145-369), N-terminal deletion of aa 19-144
CipA E360A FP	GTGTTAGATATGGTTGTAGCGCAAGA ACTCGTTAAAAAGCC	Generation of variant CipA _{E360A}
CipA E360A RP	CTTTTAACGAGTTCTGCGCTACAAC CATATCTAACACATC	Generation of variant CipA _{E360A}
CipA E360P_II FP	GATGTGTAGATATGGTTGTACCGCA AGAACTCGTTAAAAAGCC	Generation of variant CipA _{E360P}
CipA E360P_II RP	CTTTTAACGAGTTCTGCGGTACAAC CATATCTAACACATCAGC	Generation of variant CipA _{E360P}
CipA Q361A FP	GTGTTAGATATGGTTGTAGAGGCAGA ACTCGTTAAAAAGCC	Generation of variant CipA _{Q361A}
CipA Q361A RP	CTTTTAACGAGTTCTGCCCTACAAC CATATCTAACACATC	Generation of variant CipA _{Q361A}
CipA E362A FP	GATATGGTTGTAGAGCAAGCAACTCG TTAAAAAGCCTAAAAAA	Generation of variant CipA _{E362A}
CipA E362A RP	GGCTTTTAACGAGTGCTTGCTCTACA ACCATATCTAACAC	Generation of variant CipA _{E362A}
CipA P367A FP	CAAGAACTCGTTAAAAGGCTAAAAAA ATAAGTCGACCTGC	Generation of variant CipA _{P367A}
CipA P367A RP	GCTTGGCTGCAGGTCGACTTATTTC AGCCTTTAACGAG	Generation of variant CipA _{P367A}
CipA delta L363 FP	GGTTGTAGAGCAAGAAGTTAAAAGC CTAAAAAAATAAGTC	Generation of variant CipA _{ΔL363} lacking a leucine at aa position 363
CipA delta L363A RP	AGGCTTTAACCTCTGCTCTACAAC CATATCTAACAC	Generation of variant CipA _{ΔL363} lacking a leucine at aa position 363
CipA delta V364 FP	GTAGAGCAAGAACTCAAAAAGCCTA AAAAATAAGTCGACCTG	Generation of variant CipA _{ΔV364} lacking a valine at aa position 364
CipA delta V364 RP	GAGTTATTTTAGGCTTTGAGTT TTGCTCTACAAACC	Generation of variant CipA _{ΔV364} lacking a valine at aa position 364
CipA delta P367 FP	CAAGAACTCTTAAAAGAAAAATA AGTCGACCTGC	Generation of variant CipA _{ΔP367} lacking a proline at aa position 367
CipA delta P367 RP	GCTTGGCTGCAGGTCGACTTATTTC CTTTAACGAG	Generation of variant CipA _{ΔV367} lacking a proline at aa position 367
CipA delta L363-V364 FP	GTTGTAGAGCAAGAAAAAAAGCCTA AAAAATAAGTCGACCTG	Generation of variant CipA _{ΔL363-V364} lacking leucine and valine at positon 363 and 364
CipA delta L363-V364 RP	GAGTTATTTTAGGCTTTCTTG CTCTACAAACC	Generation of variant CipA _{ΔL363-V364} lacking leucine and valine at positon 363 and 364
CipA-364 FP	GTAGAGCAAGAACTCGTTAATAACC AAAAATAAGTCGAC	Generation of variant CipA _{ΔV364-K369} lacking aa at positions 364 to 369
CipA-364 RP	GTAGAGCAAGAACTCGTTAATAACC AAAAATAAGTCGAC	Generation of variant CipA _{ΔV364-K369} lacking aa at positions 364 to 369

CipA-L363 FP	GTTGTAGAGCAAGAACTCTAATAAAAA GCCTAAAAAATAAGTCGAC	Generation of variant CipA _{ΔL363-K369} lacking aa at positions 363 to 369
CipA-L363 RP	CGACTTATTTTAGGCTTTATTAGA GTTCTTGCTCTAC	Generation of variant CipA _{ΔL363-K369} lacking aa at positions 36 to 369
CipA-E362 FP	GTTGTAGAGCAAGAATAATAAAAAAA AGCCTAAAAAATAAGTC	Generation of variant CipA _{ΔE362-K369} lacking aa at positions 362 to 369
CipA-E362 RP	CTTATTTTAGGCTTTATTATTTC TTGCTCTAC	Generation of variant CipA _{ΔE362-K369} lacking aa at positions 362 to 369
CipA-Q361 FP	GGTTGTAGAGCAATAATAAGTTAAAA AGCCTAAAAAAT	Generation of variant CipA _{ΔQ361-K369} lacking aa at positions 361 to 369
CipA-Q361 RP	GGCTTTAACCTATTATTCCCTCTACA ACCATATC	Generation of variant CipA _{ΔQ361-K369} lacking aa at positions 361 to 369
CipA-E360 FP	GATATGGTTGTAGAGTAATAACTCGT TAAAAAGCCTAAAAAAT	Generation of variant CipA _{ΔE360-K369} lacking aa at positions 360 to 369
CipA-E360 RP	GGCTTTAACGAGTTATTACTCTACA ACCATATCTAAC	Generation of variant CipA _{ΔE360-K369} lacking aa at positions 360 to 369

Generation of *A. baumannii* 19606 strain producing a CipA variant by markerless mutagenesis

CipA up fwd PstI	GCGACT <u>GCAGCAA</u> ACTCAGGTTATTG AACTCCCAATGG	Amplification of the upstream region of the CipA encoding gene in strain <i>A. baumannii</i> 19606
CipA down rev NotI	GACAG <u>CGGCCGCGGG</u> ATTAAATATT TGCTGCTAAATG	Amplification of the downstream CipA encoding gene in strain <i>A. baumannii</i> 19606
CipA V359 RP	GTTAGATATGGTTGTATAGTAATAAC TCGTTAAAAGCCTAAAAAA	Introduction of two stop codons after position 359
CipA V359 FP	GGCTTTAACGAGTTATTACTATACA ACCATATCTAACACATC	Introduction of two stop codons after position 359
CipA control fwd	GCTCTGTCTATCTTATGTCACAGATA GCC	Control of the complementation in <i>A. baumannii</i> 19606
CipA control rev	CTGGCTGTCCACCAGGAACACATTAA TTTGC	Control of the complementation in <i>A. baumannii</i> 19606
CipA seq Fwd	CAATCCAAACCAACGCGTAATCTTAC GTG	Sequencing of <i>cipA</i> gene in the complemented <i>A. baumannii</i> 19606
CipA seq Rev	GGTACTGTGAATTATGTAGATGGTGC C	Sequencing of <i>cipA</i> gene in the complemented <i>A. baumannii</i> 19606

Primer used for RT-PCR of *A. baumannii* genes

rpoB-RT-fwd	GAGTCTAATGGCGGTGGTTC	RT-PCR, amplification of <i>rpoB</i> in <i>A. baumannii</i>
rpoB-RT-rev	ATTGCTTCATCTGCTGGTTG	RT-PCR, amplification of <i>rpoB</i> in <i>A. baumannii</i>
CipA-RT-PCR Fwd	GTCATGTCAACTATGATGGC	RT-PCR, amplification of <i>cipA</i> in <i>A. baumannii</i>
CipA-RT-PCR Rev	GCAAGCAAGTTGTGGTGCAAC	RT-PCR, amplification of <i>cipA</i> in <i>A. baumannii</i>

Primer used for the generation of His-tagged Efb and Efb-C proteins of *Staphylococcus aureus* USA300

Efb_FP Bam	GCGAG <u>CGAAGGATCCGGTCCAAGAG</u>	Amplification of Efb encoding gene of
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Efb_RP_Sal	AAAAGAAACCACTGAG GGCTGCAG <u>GGT</u> CGACTTATTAACTAA TCCTTGTTTAATAC	<i>S. aureus</i> USA300 Amplification of Efb encoding gene of <i>S. aureus</i> USA300
Efb-C_FP_Bam	GGTGCAGGAT <u>CCCA</u> ATTAAATAAACCG AGCAGCGAAA <u>ACTG</u>	Generation of a C-terminal fragment of Efb

Primer used for sequencing

pQE-FP-30	TTGCTTGTGAGCGGATAAC	Sequencing of pQE-30 Xa vector
pQE-RP	CTGAGGTCA <u>T</u> ACTGGATCTATC	Sequencing of pQE-30 Xa vector

^a, Sequences of specific restriction endonuclease recognition sites are underlined