

Supplementary information

Redundant and Distinct Roles of Secreted Protein Eap and Cell Wall-Anchored Protein SasG in Biofilm Formation and Pathogenicity of *Staphylococcus aureus*

Running Title: Roles of Eap and SasG in Biofilm and Pathogenesis

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Supplementary Methods

Purification of recombinant Eap. Recombinant Eap from MR23 was expressed and purified using a *Brevibacillus* expression system as previously described (1). Purified Eap was subjected to SDS-PAGE and quantified by measuring the band intensity using LAS-4000 Image Analyzer (GE Healthcare). Bovine serum albumin was used as a standard. Then, purified Eap was supplemented into biofilm cultures to address whether it promotes biofilm formation.

Atmospheric scanning electron microscopy (ASEM). *S. aureus* MR23 wild-type and Δeap strains were grown in BHIG for 37°C for 4 h in ASEM dishes, as previously described (2). Biofilms formed on the dishes were fixed with 1% (v/v) glutaraldehyde/1% (v/v) paraformaldehyde and labeled with positively charged Nanogold reagent (Nanoprobes, Stony Brook, NY). After gold enhancement using GoldEnhance-EM (Nanoprobes) for 10 min at 25°C, biofilms were washed three times with doubly distilled water. Subsequently, biofilms were stained overnight with 2% (w/v) phosphate tungstic acid at 4°C. After washing three times with doubly distilled water, biofilms were immersed in radical scavenger 1% (w/v) ascorbic acid solution and imaged by ASEM (3) using the ClairScope ASEM system (JASM-6200, JEOL, Ltd, Tokyo, Japan). The acceleration voltage of the inverted SEM was 30 kV, and the applied electron dose was less than $0.8 \text{ e}^-/\text{\AA}^2$.

Supplementary Figure Legends

FIG S1 Enzymatic susceptibility of preformed *S. aureus* biofilms. For the experiments, 24-h biofilms formed by WT (A), Δeap (B), $\Delta eap \Delta srtA$ (C), $\Delta sasG$ (D), and $\Delta eap \Delta sasG$ (E) strains were treated with the indicated enzymes or left untreated. Biomasses of the residual biofilms were quantified as described in Fig. 1. Means and standard deviations from three independent experiments are shown. **, $P < 0.01$; NS, not significant.

FIG S2 Characterization of SasG in various *S. aureus* strains. Expression levels of SasG were analyzed by immunoblotting for clinical strains of methicillin-resistant (MRSA) and -sensitive *S. aureus* (MSSA) that produce protein-dependent biofilms. Molecular sizes are also indicated at the left of the panels.

FIG S3 Biofilm formation by *S. aureus* RN4220 and its isogenic mutant strains. (A) Biofilms of the indicated RN4220-derived strains were formed and quantified as indicated in Figure 2A. (B) Biofilm formation by RN4220 $\Delta eap \Delta sasG$ -expressing Eap, SasG, and both was also analyzed. The optimal concentrations of anhydrotetracycline (aTc) were added to induce the expression of each protein. (C) SasG in the cell wall fraction and Eap in the ECM fraction from the strains grown under the indicated condition in B were detected by SDS-PAGE with CBB-staining. Red arrows indicate SasG. (D) Biofilm formation of RN4220 $\Delta eap \Delta sasG$ harboring pLC1 or pEap-SasG was analyzed in the presence of 3.13 ng/ml aTc. The indicated concentrations of purified Eap was added to biofilm cultures from the onset of biofilm formation. The means and standard deviations of biofilm biomasses from three independent experiments are shown. *, $P < 0.05$; **, $P < 0.01$; NS, not significant.

FIG S4 ASEM analysis of biofilms. ASEM images of 4-h biofilms formed by MR23 WT (A) and $\Delta eap \Delta sasG$ (B) strains labeled with positively charged Nanogold and phosphate tungstic acid are shown. Scale bars, 10 μm ($\times 1,000$ magnification) and 1 μm ($\times 10,000$ magnification).

FIG S5 Proliferation of *S. aureus* in silkworm larvae. Silkworm larvae were sacrificed and

body fluids collected 24 h after injection of the indicated *S. aureus* strains. The body fluids were diluted and spread on mannitol salt agar plates. The plates were incubated at 37°C overnight and the obtained colonies counted. Means and standard deviations from three independent experiments are shown. NS, not significant.

TABLE S1 Bacterial strains and plasmids used in the current study.

Strains and plasmids	Description	Source or reference
<i>S. aureus</i> strains		
RN4220	NCTC8325-4 derivative, restriction deficient mutant	4
RN4220 Δeap	The <i>eap</i> gene was deleted from RN4220	This study
RN4220 $\Delta sasG$	The <i>sasG</i> gene was deleted from RN4220	This study
RN4220 $\Delta eap \Delta sasG$	The <i>eap</i> and <i>sasG</i> genes were deleted from RN4220	This study
MR23	A clinical isolate of MRSA from the Jikei hospital	1
MR23 Δeap	The <i>eap</i> gene was deleted from MR23	This study
MR23 $\Delta srtA$	The <i>srtA</i> gene was deleted from MR23	5
MR23 $\Delta eap \Delta srtA$	The <i>eap</i> and <i>srtA</i> genes were deleted from MR23	This study
MR23 $\Delta sasG$	The <i>sasG</i> gene was deleted from MR23	This study
MR23 $\Delta eap \Delta sasG$	The <i>eap</i> and <i>sasG</i> genes were deleted from MR23	This study
MR23 $\Delta eap \Delta fnbA \Delta fnbB$	The <i>eap</i> , <i>fnbA</i> and <i>fnbB</i> genes were deleted from MR23	This study
MR23 $\Delta eap \Delta clfA \Delta clfB$	The <i>eap</i> , <i>clfA</i> and <i>clfB</i> genes were deleted from MR23	This study
MR23 $\Delta eap \Delta spa$	The <i>eap</i> and <i>spa</i> genes were deleted from MR23	This study
MR23 $\Delta eap \Delta isdAB$	The <i>eap</i> and <i>isdAB</i> genes were deleted from MR23	This study
<i>E. coli</i> strain		
DH5 α	<i>fhuA2</i> Δ (<i>argF-lacZ</i>) <i>U169 phoA glnV44 ϕ80</i> Δ (<i>lacZ</i>) <i>M15 gyrA96 recA1 relA1 endA1 thi-1 hsdR17</i>	Toyobo, Osaka, Japan
BLR (DE3)	<i>F⁻ ompT hsdS_B(r_B⁻ m_B⁻) gal lac ile dcm</i> Δ (<i>srl-recA</i>) <i>306::Tn10 (tet^R)(DE3)</i>	Novagen
Plasmids		
pKOR1	<i>E. coli-S. aureus</i> shuttle vector plasmid for knockout of genes by allelic exchange, Amp ^R , Cm ^R	6
pSrtA-ko	pKOR1-derivative plasmid for knockout of the MR23 <i>srtA</i> gene, Amp ^R , Cm ^R	5
pEap-ko	pKOR1-derivative plasmid for knockout of the MR23 <i>eap</i> gene, Amp ^R , Cm ^R	This study
pSasG-ko	pKOR1-derivative plasmid for knockout of the MR23 <i>sasG</i> gene, Amp ^R , Cm ^R	This study
pFnbA-ko	pKOR1-derivative plasmid for knockout of the MR23 <i>fnbA</i> gene, Amp ^R , Cm ^R	This study
pFnbB-ko	pKOR1-derivative plasmid for knockout of the MR23 <i>fnbB</i> gene, Amp ^R , Cm ^R	This study
pClfA-ko	pKOR1-derivative plasmid for knockout of the MR23 <i>clfA</i> gene, Amp ^R , Cm ^R	This study
pClfB-ko	pKOR1-derivative plasmid for knock out of the MR23 <i>clfB</i> gene, Amp ^R , Cm ^R	This study
pSpa-ko	pKOR1-derivative plasmid for knock out of the MR23 <i>spa</i> gene, Amp ^R , Cm ^R	This study
pIsdAB-ko	pKOR1-derivative plasmid for knock out of the MR23 <i>isdAB</i> gene, Amp ^R , Cm ^R	This study
pLC1	<i>E. coli-S. aureus</i> shuttle vector plasmid with <i>tetO/xyl</i> promoter, Amp ^R , Cm ^R	7
pEap	pLC1-derivative plasmid for expression of Eap, Amp ^R , Cm ^R	This study
pSrtA	pLC1-derivative plasmid for expression of Sortase A, Amp ^R , Cm ^R	This study
pSasG ^{WT}	pLC1-derivative plasmid for expression of SasG ^{WT} , Amp ^R , Cm ^R	This study
pSasG ^{ΔL}	pLC1-derivative plasmid for expression of SasG ^{ΔL} , Amp ^R , Cm ^R	This study
pEap-SasG ^{WT}	pLC1-derivative plasmid for co-expression of Eap and SasG ^{WT} , Amp ^R , Cm ^R	This study

pCold I	Cold shock promoter, His ₆ -tag gene, ColE1 <i>ori</i> , Ap ^R	Takara
pCold-SasG	pCold I-derivative plasmid for expression of SasG(51-954)	This study

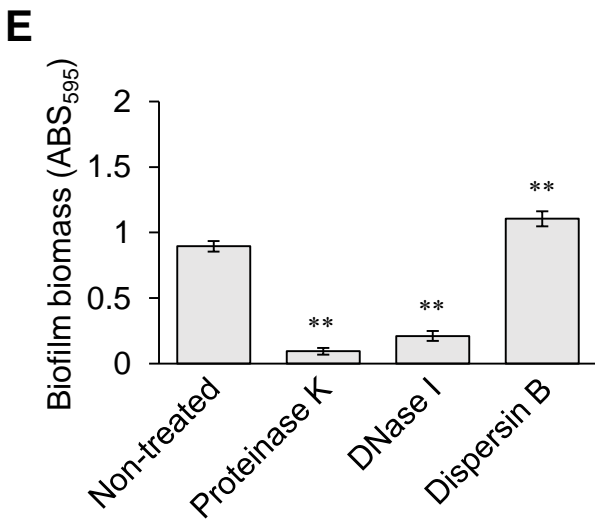
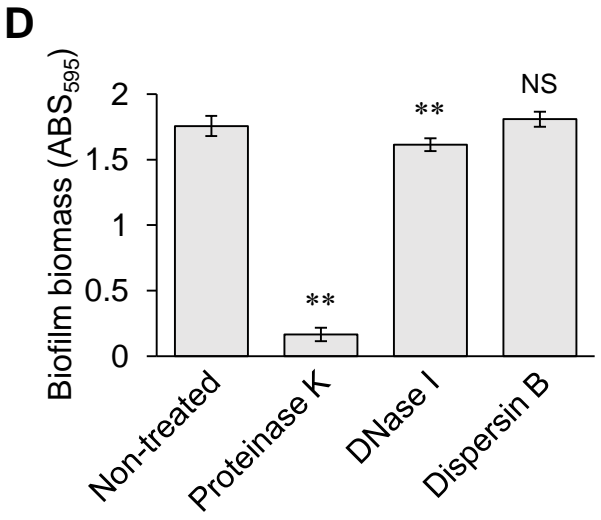
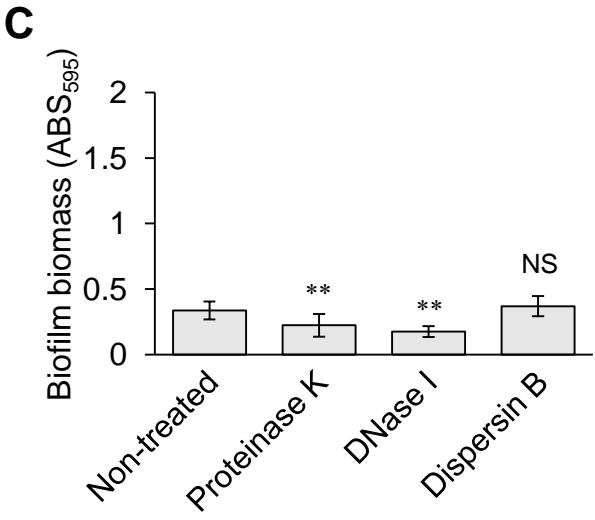
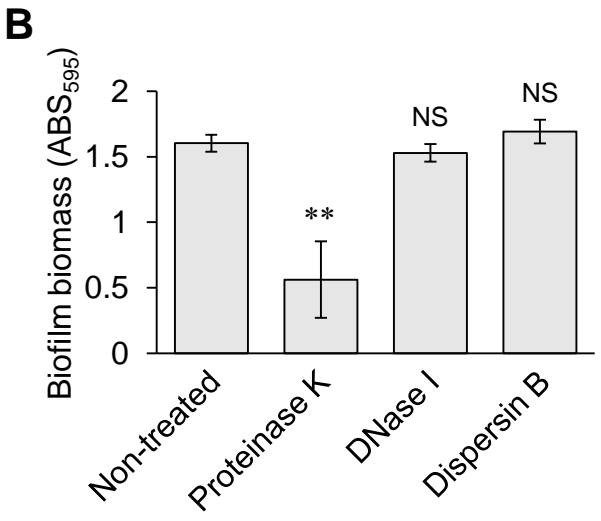
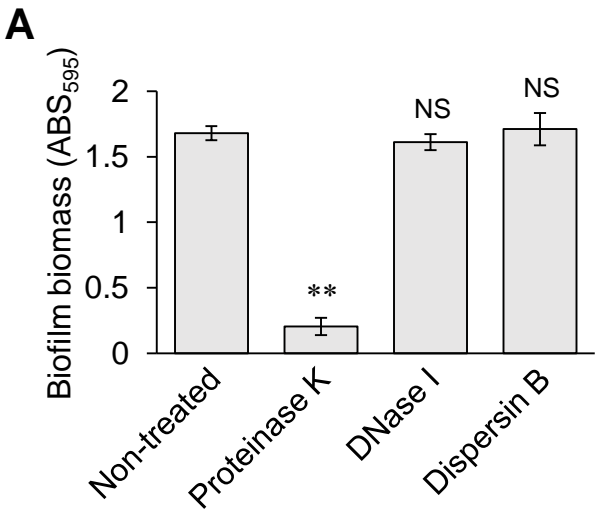
Amp^R, ampicillin-resistant; Cm^R, chloramphenicol-resistant; tet^R, tetracycline-resistant .

TABLE S2 Oligonucleotide primers used in the current study.

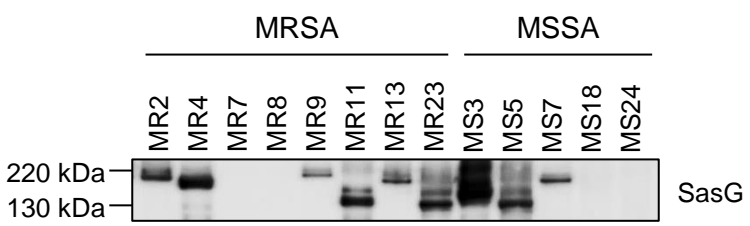
Primers	Sequences (5' to 3')	Descriptions
attB1- <i>eap</i> -F	ggggacaagttgtacaaaaaagcaggctataaaagttcatgatgttgacc	Disruption of <i>eap</i> in MR23
<i>eap</i> -R	gaattacacaaaaaaggagagataattttcttaattttataataaggcatctcac	Disruption of <i>eap</i> in MR23
<i>eap</i> -F	gtgagatgccttattataaaaattaaagaaaattatctctcctttttgtgaattc	Disruption of <i>eap</i> in MR23
attB2- <i>eap</i> -R	ggggaccactttgtacaagaaagctgggttcagtaagtcaccaactaataataat	Disruption of <i>eap</i> in MR23
attB1- <i>srtA</i> -F	ggggacaagttgtacaaaaaagcaggcttaataatctattttcactcgttatctta	Disruption of <i>srtA</i> in MR23
<i>srtA</i> -R	atccatccattagcgtaatagaacgtaaggctcctttataca	Disruption of <i>srtA</i> in MR23
<i>srtA</i> -F	tgtataaaaggagccttaacgcttattacgctaattggatgaat	Disruption of <i>srtA</i> in MR23
attB2- <i>srtA</i> -R	ggggaccactttgtacaagaaagctgggttaaccatctattaaatataaacctacatt	Disruption of <i>srtA</i> in MR23
attB1- <i>sasG</i> -F	ggggacaagttgtacaaaaaagcaggctagattagcatatttagatgtattaaaaa	Disruption of <i>sasG</i> in MR23
<i>sasG</i> -R	aatcaactattatataaattatgaatttttgcatactccttttcca	Disruption of <i>sasG</i> in MR23
<i>sasG</i> -F	tggaaaaaggagatgcaaaaataatcataatttaataatagttgatt	Disruption of <i>sasG</i> in MR23
attB2- <i>sasG</i> -R	ggggaccactttgtacaagaaagctgggtcatcattgataatttaaagatgat	Disruption of <i>sasG</i> in MR23
attB1- <i>fnbA</i> -F	ggggacaagttgtacaaaaaagcaggctcgttcaggatcatctcattttattc	Disruption of <i>fnbA</i> in MR23
<i>fnbA</i> -R	gaaataaacccgtcaattttgttaataataatctcctttaaattgcaaaattca	Disruption of <i>fnbA</i> in MR23
<i>fnbA</i> -F	tgaattttgcattaaaggagatattatataaacaaaaattgacgggtttttc	Disruption of <i>fnbA</i> in MR23
attB2- <i>fnbA</i> -R	ggggaccactttgtacaagaaagctgggtcaataaatgcaattctatattgttcgg	Disruption of <i>fnbA</i> in MR23
attB1- <i>fnbB</i> -F	ggggacaagttgtacaaaaaagcaggctcaaagcataattaaacaaaaattgacg	Disruption of <i>fnbB</i> in MR23
<i>fnbB</i> -R	aaataaacctgtcaattttggattgatataatattctccttaaattgcaaaattca	Disruption of <i>fnbB</i> in MR23
<i>fnbB</i> -F	tgaattttgcattaaaggagaatattatcaatccaaaattgacagggtttttt	Disruption of <i>fnbB</i> in MR23
attB2- <i>fnbB</i> -R	ggggaccactttgtacaagaaagctgggtgataaattgacgcgcagaactaat	Disruption of <i>fnbB</i> in MR23
attB1- <i>clfA</i> -F	ggggacaagttgtacaaaaaagcaggctttgatattcatcatcaaaataaa	Disruption of <i>clfA</i> in MR23
<i>clfA</i> -R	atcatatgattaatataatcattattactttattccctcttttaaaaaag	Disruption of <i>clfA</i> in MR23
<i>clfA</i> -F	cttttaaaaaagaggggaataaagtaataatgatataaattaatcatatgat	Disruption of <i>clfA</i> in MR23
attB2- <i>clfA</i> -R	ggggaccactttgtacaagaaagctgggttatttttctctcactaaaaaat	Disruption of <i>clfA</i> in MR23
attB1- <i>clfB</i> -F	ggggacaagttgtacaaaaaagcaggcttgactgtcaaattttgatgcata	Disruption of <i>clfB</i> in MR23
<i>clfB</i> -R	taatctagaaattgaaatggagtaatatttagactcggatagcgaactcag	Disruption of <i>clfB</i> in MR23
<i>clfB</i> -F	ctgagtcgctatccgagctaaatattactccatttcaatttctagatta	Disruption of <i>clfB</i> in MR23
attB2- <i>clfB</i> -R	ggggaccactttgtacaagaaagctgggtatgaattttatgaaatcaaacagttttt	Disruption of <i>clfB</i> in MR23
attB1- <i>isdAB</i> -F	ggggacaagttgtacaaaaaagcaggctacagttatttgaacatacaatttacc	Disruption of <i>isdAB</i> in MR23
<i>isdAB</i> -R	aatttaataatataaataaaagacgatttagttgtttcctcctaaggata	Disruption of <i>isdAB</i> in MR23
<i>isdAB</i> -F	tatccttaggaggaaaacaactaaatcgtcttatattaattattaaatt	Disruption of <i>isdAB</i> in MR23
attB2- <i>isdAB</i> -R	ggggaccactttgtacaagaaagctgggtgaagtaatcgaatgaggatcaaT	Disruption of <i>isdAB</i> in MR23
attB1- <i>spa</i> -F	ggggacaagttgtacaaaaaagcaggctgaacgctcaactgaagatgaaag	Disruption of <i>spa</i> in MR23
<i>spa</i> -R	gatatctatcgttgattgtttgttaataataacgaattatgtattgcaatac	Disruption of <i>spa</i> in MR23
<i>spa</i> -F	gtattgcaatacataattcgttatattaacaaacaatacacacgatagatc	Disruption of <i>spa</i> in MR23
attB2- <i>spa</i> -R	ggggaccactttgtacaagaaagctgggtcattactgtggcagctaacac	Disruption of <i>spa</i> in MR23
pLC1-F	ctgcagccaagctagcttggc	Linearization of pLC1 vector
pLC1-R	cccgggagatctgatatcaagcttatttta	Linearization of pLC1 vector

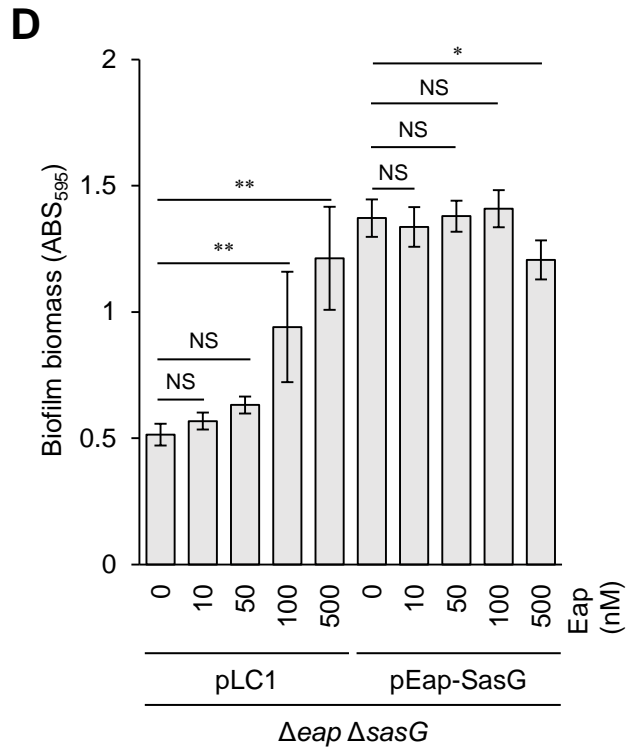
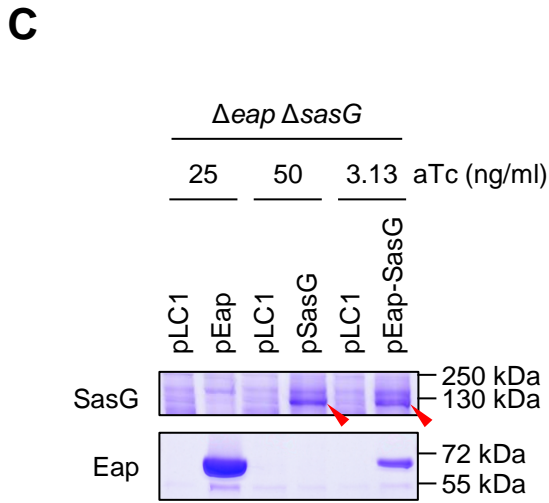
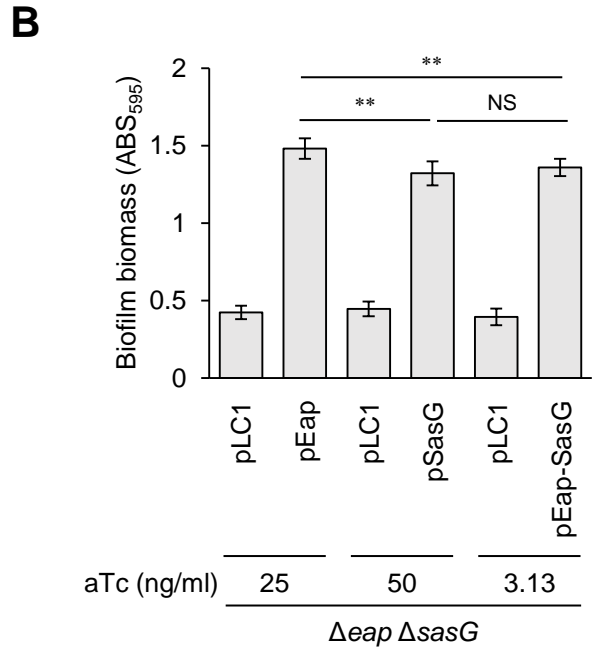
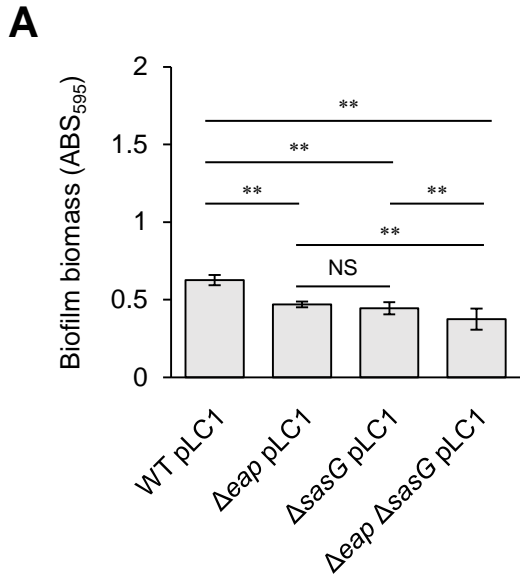
pEap-F	atcagatctcccggggcttggtataaaacaaaaccaaaca	Construction of Eap-expressing plasmid
pEap-R	ctagcttgctgcagatcattaattgtaattttgcttgattata	Construction of Eap-expressing plasmid
pSrtA-F	atcagatctcccgggtgtataaaaggagcctaacgtatga	Construction of SrtA-expressing plasmid
pSrtA-R	ctagcttgctgcagttattgacttctgtagctacaaagatttt	Construction of SrtA-expressing plasmid
pSasG ^{WT} -F	atcagatctcccgggtggaaaaaggagatgcaaag	Construction of SasG ^{WT} -expressing plasmid
pSasG ^{WT} -R	ctagcttgctgcagtaattctttctctacgagccaata	Construction of SasG ^{WT} -expressing plasmid
pSasG ^{ΔL} -F	taactgcagccaagctagcttggc	Construction of SasG ^{ΔL} -expressing plasmid
pSasG ^{ΔL} -R	gcttggctgcagttattctgctcgtttttctcttgat	Construction of SasG ^{ΔL} -expressing plasmid
pEap-SasG ^{WT} -F1	ctgcagccaagctagcttggc	Construction of Eap-SasG ^{WT} -co-expressing plasmid
pEap-SasG ^{WT} -R1	atcattaattgtaattttgcttgattatagaacacaa	Construction of Eap-SasG ^{WT} -co-expressing plasmid
pEap-SasG ^{WT} -F2	attacaattaatgattggaaaaaggagatgcaaag	Construction of Eap-SasG ^{WT} -co-expressing plasmid
pEap-SasG ^{WT} -R2	ctagcttgctgcagtaattctttctctacgagccaata	Construction of Eap-SasG ^{WT} -co-expressing plasmid
pCold-SasG-F	gaaggtaggcatatggctgaaaaacaatattgagaatccaactacattaaaagataatgtcc	Construction of SasG(51-954)-expressing plasmid
pCold-SasG-R	agagattacctatctttattctgctcgtttttctcttgattagctactgattctttagc	Construction of SasG(51-954)-expressing plasmid
SasG-s1	agtattagcatatttagatgtatttaaaaa	For DNA sequencing of sasG in MR23
SasG-s2	tagttgaatctacccaattacaattcaag	For DNA sequencing of sasG in MR23
SasG-s3	tatgCGGacaattcaactaatacatcagat	For DNA sequencing of sasG in MR23
SasG-s4	ccaaaattacaacctgggaagagcgagtg	For DNA sequencing of sasG in MR23
SasG-s5	gtaaattcactgtaagtaaagtggaaaata	For DNA sequencing of sasG in MR23
SasG-s6	tcgaaagaagaaatcacaaaagatccgatt	For DNA sequencing of sasG in MR23
SasG-s7	acaaaagaagagattacaaaagatccgat	For DNA sequencing of sasG in MR23

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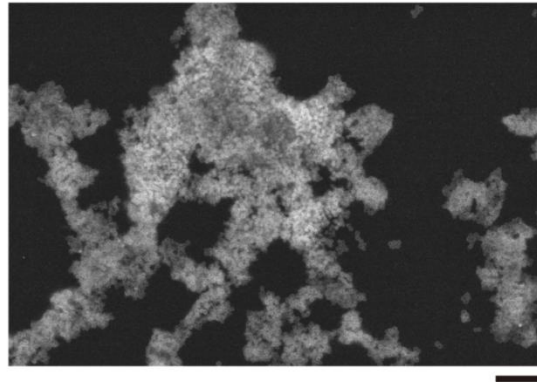
Yonemoto *et al.* Figure S2



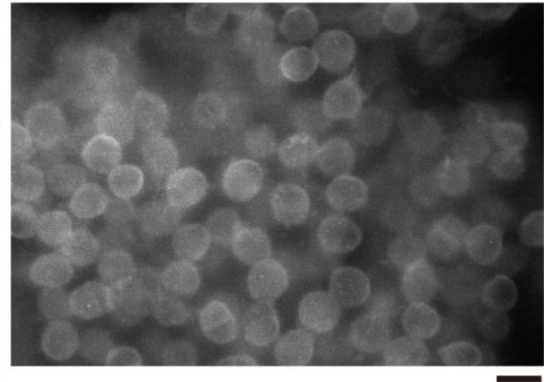


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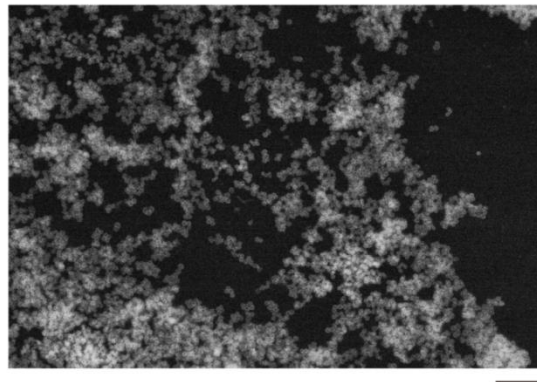


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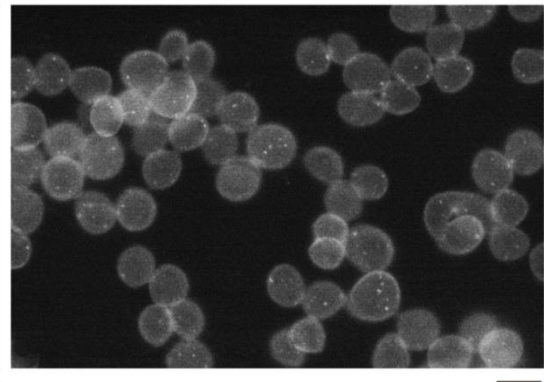


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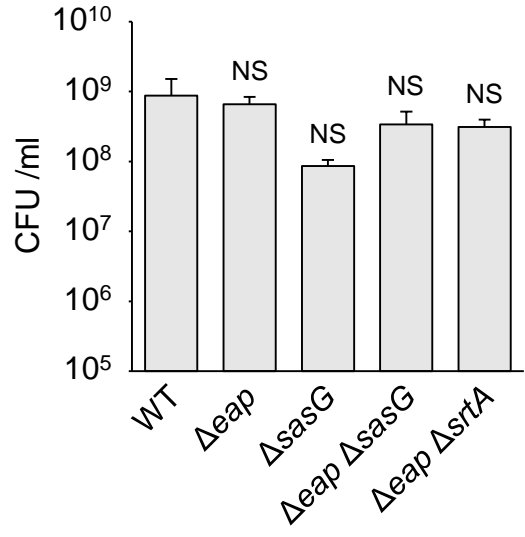
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Yonemoto *et al.* Figure S5



Supplementary References

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