A Novel Bacterial Resistance Mechanism against Human Group IIA-Secreted Phospholipase A₂: Role of *Streptococcus pyogenes* Sortase A

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A Novel Bacterial Resistance Mechanism Against Human Group IIA-Secreted Phospholipase A2: Role of \textit{Streptococcus pyogenes} Sortase A

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Human group IIA-secreted phospholipase A2 (sPLA₂-IIA) is a bactericidal molecule important for the innate immune defense against Gram-positive bacteria. In this study, we analyzed its role in the host defense against \textit{Streptococcus pyogenes}, a major human pathogen, and demonstrated that this bacterium has evolved a previously unidentified mechanism to resist killing by sPLA₂-IIA. Analysis of a set of clinical isolates demonstrated that an ∼500-fold higher concentration of sPLA₂-IIA was required to kill \textit{S. pyogenes} compared with strains of the group B \textit{Streptococcus}, which previously were shown to be sensitive to sPLA₂-IIA, indicating that \textit{S. pyogenes} exhibits a high degree of resistance to sPLA₂-IIA. We found that an \textit{S. pyogenes} mutant lacking sortase A, a transpeptidase responsible for anchoring LPXTG proteins to the cell wall in Gram-positive bacteria, was significantly more sensitive (∼30-fold) to sPLA₂-IIA compared with the parental strain, indicating that one or more LPXTG surface proteins protect \textit{S. pyogenes} against sPLA₂-IIA. Importantly, using transgenic mice expressing human sPLA₂-IIA, we showed that the sortase A-mediated sPLA₂-IIA resistance mechanism in \textit{S. pyogenes} also occurs in vivo. Moreover, in this mouse model, we also showed that human sPLA₂-IIA is important for the defense against lethal \textit{S. pyogenes} infection. Thus, we demonstrated a novel mechanism by which a pathogenic bacterium can evade the bactericidal action of sPLA₂-IIA and we showed that sPLA₂-IIA contributes to the host defense against \textit{S. pyogenes} infection. \textit{The Journal of Immunology}, 2011, 187: 000–000.
lated LTA (23), the sensitivity to sPLA2-IIA differs greatly between species (11, 12), indicating that some Gram-positive pathogens may have evolved other resistance mechanisms to sPLA2-IIA.

In this study, we identified a novel bacterial molecule, the transpeptidase sortase A (SrtA) of *Streptococcus pyogenes*, that is involved in the resistance to sPLA2-IIA. *S. pyogenes* is a major Gram-positive pathogen causing a wide range of clinical manifestations in humans, such as pharyngitis, skin infections, and a toxic shock syndrome. It also gives rise to two important post-infectious sequelae: acute rheumatic fever and acute glomerulonephritis (24). *S. pyogenes* expresses several cell surface proteins that are important for various steps in the molecular pathogenesis of *S. pyogenes* infection, such as phagocytosis resistance, adhesion, and invasion of host cells (25, 26). Many of these cell surface proteins, including the antiphagocytic M protein, belong to a class of proteins containing a C-terminal LPXTG motif that are anchored to the bacterial cell wall peptidoglycan through a process that involves the transpeptidase SrtA (27, 28). SrtA recognizes the LPXTG motif and catalyzes the transpeptidation reaction that results in covalent binding of the protein to peptides in cell wall peptidoglycan (27, 28). Studies on Gram-positive pathogens showed that SrtA− mutants are significantly less virulent compared with their wild-type (WT) counterparts in animal models (29–32), demonstrating that LPXTG proteins, in general, are important virulence factors.

In this article, we show that clinical strains of *S. pyogenes* exhibit a high degree of resistance against sPLA2-IIA. In an attempt to identify the bacterial factors that contribute to this resistance, we found that SrtA is important for *S. pyogenes* to resist direct killing by sPLA2-IIA, indicating that one or more LPXTG surface proteins are involved in this innate immune-evasion mechanism. Moreover, using Tg mice expressing human sPLA2-IIA, we present evidence that this resistance mechanism also occurs in vivo and that sPLA2-IIA contributes to the innate immune defense against lethal *S. pyogenes* infection. Thus, we demonstrate a novel mechanism by which a pathogenic bacterium can evade the direct bactericidal action of sPLA2-IIA and, to our knowledge, we show for the first time that sPLA2-IIA contributes to the host defense against *S. pyogenes* infection.

### Materials and Methods

#### Bacterial strains

The WT *S. pyogenes* serotype M6 strain JRS4, the isogenic SrtA− mutant strain (JRS758), and the sortase B (SrtB)-mutant strain (JRS789) were constructed as described (33) and kindly provided by Dr. J.R. Scott (Emory University School of Medicine, Atlanta, GA). Plasmid pJRS757, containing the intact srtA gene, was constructed as described (33) and used for trans-complementation of the SrtA− mutant. A collection of clinical *S. pyogenes* strains representing serotypes Ia, Iii, III, III, and V, respectively, were kindly provided by Dr. G. Lindahl (Lund University). Strains of *S. pyogenes* were grown in Todd–Hewitt broth supplemented with 0.2% yeast extract (THY) in 5% CO₂ at 37°C and GBS strains were grown in Todd–Hewitt broth at 37°C.

#### Proteins and reagents

Recombinant WT human sPLA2-IIA was produced in *Drosophila* S2 cells using a protocol similar to that previously used for production in baculovirus-infected insect cells (34) and following the use of insect cells for the structurally related human group IID sPLA2 (35). The sPLA2-IIA preparation was checked for purity by SDS-PAGE analysis, MALDI-TOF mass spectrometry, and sPLA2-IIA enzymatic assays (35). Before use, the purified lipophylic sPLA2-IIA preparation was solubilized in PBS. The sPLA2-IIA-specific inhibitor LY311727 was described previously (36). The human AMP LL-37 and Magainin II were purchased from Sigma (St. Louis, MO).

#### Bactericidal analysis with recombinant human sPLA2-IIA and AMPs

The sPLA2-IIA bactericidal assay was adapted from a previously described protocol (8, 9). Briefly, overnight cultures of streptococci (OD₆₀₀ = 1.1) were diluted 20-fold into fresh broth, and bacteria were grown at 37°C to mid-log phase (OD₆₀₀ = 0.4). Bacteria were then directly diluted 1000-fold into HEPES buffer (20 mM HEPES, 2 mM Ca²⁺, 1% BSA [pH 7.4]) to give a bacterial concentration of ∼1 × 10⁸ CFU/ml. The recombinant human sPLA2-IIA or AMP was serially diluted in HEPES buffer, and small aliquots of 10 μl were added in triplicates to the wells of a sterile 96-well polystyrene plate (Corning), after which it was mixed with 10 μl of the bacterial suspension. After incubation at 37°C with 5% CO₂ for 2 h, the samples were serially diluted in PBS, plated onto blood agar plates, and incubated overnight to determine the bacterial viability by colony counting. In the inhibition experiment, LY311727 was added at a final concentration of 50 μM. In the analysis of the role of bacterial growth phase on sensitivity to sPLA2-IIA, bacteria were used at mid-log phase (OD₆₀₀ = 0.4) and early stationary phase (OD₆₀₀ = 0.9).

#### Radiolabeling of *S. pyogenes* membrane and analysis of bacterial phospholipid degradation

Bacterial phospholipids were labeled, essentially as described (8). Briefly, overnight cultures of streptococci were inoculated into fresh THY containing 1 μCi [³H]oleic acid ([³H]-OA) and grown to mid-log phase (OD₆₀₀ = 0.4). After centrifugation, the bacterial pellet was resuspended in fresh THY and incubated for an additional 30 min at 37°C. After washing three times in 0.15 M NaCl to remove free oleic acid, the bacteria were resuspended in HEPES buffer (20 mM [pH 7.4]) containing 1% fatty acid-free BSA and used promptly. The bacteria were incubated with 2.5 μg/ml sPLA2-IIA for 2 h at 37°C. In the inhibition experiment using LY311727, the sPLA2-IIA dilution was preincubated with 50 μM LY311727 for 30 min at 37°C before incubation with bacteria. Bacterial phospholipid degradation products generated by sPLA2-IIA treatment were quantitatively recovered from centrifuged extracellular medium as complexes with BSA and measured by liquid scintillation spectroscopy. Photolipid degradation was calculated as a percentage of total sample radioactivity.

#### sPLA2-IIA assays

Tissue homogenates were prepared, as previously described (6). Frozen tissues were suspended in 10 volumes of a lysis buffer containing 0.25 mM sucrose, 10 mM Tris-HCl ([pH 7.4]), 1 mM EDTA, 0.5 mM PMSF, 2 μg/ml leupeptin, and 2 μg/ml aprotinin and disrupted with an Ultra-Turrax T-25 (Janke and Kunkel) at 4°C. The homogenates were centrifuged for 5 min at 1,000 × g, the soluble fractions were collected and centrifuged for 20 min at 20,000 × g, and the resulting supernatant was used to evaluate sPLA2-IIA activity. sPLA2-IIA activity was assessed using [³H]oleate-labeled membranes of *E. coli*, following a modification of the method of Franson et al. (39). The identity and level of the sPLA2-IIA protein in tissue homogenates and body fluids from Tg sPLA2-IIA and corresponding littermate mice were determined by inhibition of sPLA2-IIA activity with the inhibitor LY311727, as previously described (17).

#### sPLA2-IIA–binding assays and analysis of bactericidal activity after incubation with bacteria

The binding assay with whole washed bacteria and [¹²⁵I]labeled sPLA2-IIA was modified from a previously described method (40). Mid-log–phase bacteria were washed in HEPES buffer and suspended to a 5% bacterial suspension (∼5 × 10⁷ CFU/ml). Bacteria were serially diluted and added to [¹²⁵I]labeled sPLA2-IIA (∼10,000 cpm). After incubation at room temperature (RT) for 10 min, bacteria were washed twice with PBS containing 0.02% NaN₃ and 0.01% Tween-20 (PBSAT), and radioactivity associated with the

### Mice

Female human sPLA2-IIA Tg mice (C57BL/6N-Taq2-Tgf[N(sPLA2)] and C57BL/6 WT mice were maintained, as described (37). The WT C57BL/6 mice have a natural mutation in the gene encoding mouse sPLA2-IIA ([Pla2g2a]; therefore, they are deficient for the mouse enzyme (38). In the Tg mouse model, the transgene fragment contains the entire human sPLA2-IIA gene together with a 1.6-kb region upstream of the sPLA2-IIA gene, with the promoter and response elements for diverse effectors (IL-6, IFN, hepatocyte NF-3, AP1, AP2, C/EBP, and cAMP response element) (37). All animal experiments were performed in strict accordance with guidelines from the Lund/Malmö ethical review board on animal experiments.
bacterial pellet was measured with a gamma counter. In another binding assay, mid-log–phase bacteria were washed in HEPES buffer and sus-
pended to a 1% suspension ($\sim 1 \times 10^9$ CFU/ml). Bacteria were mixed with various amounts of $[^{125}\text{I}]\text{sPLA2-IIA}$ (50,000, 10,000 and 2,000 cpm) and incubated at 37°C for 10 min. The bacteria were washed twice in PBSAT, boiled in sample buffer for 5 min, and subjected to SDS-PAGE, after which proteins were blotted onto a nitrocellulose membrane and analyzed by autoradiography.

To verify that sPLA2-IIA was bound to the surface of bacteria, mid-log–phase JRS4 bacteria were washed once in HEPES buffer and incubated with 100 $\mu$M sPLA2-IIA (final concentration, 12.5 $\mu$g/ml) for 10 min at 37°C. The bacteria were centrifuged, and the supernatant (containing sPLA2) was diluted $\times 100$, $\times 100$, and $\times 500$ in HEPES buffer, and a killing assay analyzing the bactericidal activity in the eluates against GBS strain BM110 (grown to mid-log phase) was performed. Aliquots of diluted eluates were added in triplicates to a sterile 96-well polypropylene plate and mixed with the bacterial suspension ($\sim 1 \times 10^9$ CFU/ml). After incubation at 37°C for 30 min, the samples were diluted in PBS and plated onto blood agar plates to determine the bacterial viability by colony counting. The inhibitor LY311727 (50 $m$) was used in control wells to verify that the bactericidal activity was specific for sPLA2-IIA in the eluates.

**Mouse experiments**

Groups of WT C57BL/6 or Tg C57BL/6N-TgN (human sPLA2-IIA) mice were infected i.p. with a 1:1 mixture of mid-log–phase JRS4 WT and SrtA mutant bacteria. This mixture was defined as the input. Each mouse received $\sim 5 \times 10^4$ CFU. Surviving mice were sacrificed after 12 h, and spleens and livers were collected. Organs from a total of 11 WT C57BL/6 and 10 Tg C57BL/6N-TgN (human sPLA2-IIA) mice were analyzed. The organs were homogenized, serially diluted in PBS, and plated on blood agar. The bacteria in these samples, defined as the output, were analyzed for the presence of the two strains in the input. For each organ, a total of 100–200 colonies were picked and grown in short patches on THY agar containing 100 $\mu$g/ml spectinomycin. On these plates, only the srtA mutant strain grows because it contains a spectinomycin resistance gene not present in the WT JRS4 strain. The competitive index (CI) was defined as the ratio between the JRS4 and SrtA strains in the output divided by the ratio in the input (41).

To analyze the role of sPLA2-IIA in the protection against lethal S. pyogenes infection, the WT strain JRS4 was grown to mid-log phase and diluted in ice-cold THY. Groups of six C57BL/6 and Tg C57BL/6N-TgN (sPLA2-IIA) mice were infected i.p. with $\sim 10^5$ CFU of JRS4 WT bacteria, representing an $\sim 90\%$ lethal dose in the C57BL/6 mice. Mice were checked every 4 h, and deaths were recorded daily.

**Statistical analysis**

CIs derived from mixed-infection experiments were analyzed with the Student t test. The significance in difference in lethality between WT C57BL/6 and Tg sPLA2-IIA mice when infected with S. pyogenes strain JRS4 was calculated with the two-tailed Fischer exact test.

**Results**

**Clinical strains of S. pyogenes show a high degree of resistance against the bactericidal activity of human sPLA2-IIA**

We analyzed bactericidal activity of human sPLA2-IIA to S. pyogenes in comparison with GBS, a major neonatal pathogen (42) previously shown to be relatively sensitive to direct killing by sPLA2-IIA (12). We selected a number of clinical S. pyogenes isolates representing several important M serotypes (M1, M4, M5, M6, M18, M22, and M28) and GBS strains representing the three clinically most important capsular serotypes (Ia, III, and V). As shown in Fig. 1, this analysis demonstrated striking differences between S. pyogenes and GBS with regard to sensitivity to the bactericidal activity of human sPLA2-IIA. For GBS, the concentration of sPLA2-IIA killing 50% of the bacteria ranged from $\sim 0.5$ to $\sim 4$ ng/ml, whereas it ranged from $\sim 0.1$ to $\sim 2$ $\mu$g/ml for S. pyogenes. Thus, on average, a $\sim 500$-fold higher concentration of sPLA2-IIA is required to kill S. pyogenes compared with GBS, demonstrating that strains of S. pyogenes exhibit a high degree of resistance to sPLA2-IIA.

S. pyogenes SrtA contributes to the resistance against human sPLA2-IIA

As described above, $\alpha$-alanylation of surface LTA was described as a major resistance mechanism to human sPLA2-IIA in Gram-

positive bacteria (18, 20). However, because both S. pyogenes and GBS express $\alpha$-alanylated LTA (43, 44), we hypothesized that strains of S. pyogenes may have developed other mechanisms to resist direct killing by sPLA2-IIA, and we focused on investigating the role of surface proteins in this context. To investigate this hypothesis, we analyzed two isogenic S. pyogenes mutants, lacking either of the sortases SrtA or SrtB, derived from the M6 strain JRS4, in our bactericidal assay. SrtA was shown to be responsible for the covalent surface attachment of several well-known LPXTG proteins in S. pyogenes implicated as virulence factors, such as the M protein, the Csa peptidase, protein F, and protein GRAB (33), whereas SrtB was shown to be responsible for cell wall anchoring of the T6 protein, which is a subunit of pili in JRS4 (33). As demonstrated in Fig. 2A, the SrtA mutant, but not the SrtB mutant, is significantly more sensitive to the bactericidal activity of human sPLA2-IIA compared with JRS4. Indeed, a $\sim 30$-fold lower concentration of sPLA2-IIA was enough to kill 50% of the SrtA mutant compared with the WT strain JRS4. To exclude that this difference is due to a polar effect or secondary mutation in the isogenic SrtA mutant, we performed trans-complementation analysis by transforming the SrtA mutant with plasmid pJR757, which contains the intact srtA gene. Trans-complementation of SrtA with pJR757 was previously demon-

strated to restore the WT phenotype in the SrtA mutant (33). As shown in Fig. 2B, the trans-complemented strain SrtA*/ pJR757 was equally resistant to sPLA2-IIA as the WT strain JRS4, showing that the sensitivity of the SrtA mutant to sPLA2-IIA is completely due to the lack of srtA. This result indicated that one or more LPXTG surface proteins contribute to the protection against sPLA2-IIA attack in S. pyogenes, demonstrating a novel bacterial-resistance mechanism against this bactericidal molecule.

Importantly, killing of the SrtA mutant was mediated by sPLA2-IIA alone, because the bactericidal activity was strongly inhibited by the sPLA2-IIA inhibitor LY311727 (Fig. 2C). In an attempt to identify the LPXTG protein in S. pyogenes involved in the resistance to sPLA2-IIA, we analyzed the sensitivity of isogenic S. pyogenes mutants lacking the M6 protein and the fibronectin-binding protein F, but they were equally as resistant to sPLA2-IIA as the parental strain JRS4 (data not shown).

To analyze the killing kinetics, we chose a concentration of sPLA2-IIA (2.5 $\mu$g/ml; Fig. 2A) at which the SrtA mutant bacteria are completely killed and the WT JRS4 bacteria are only partially killed after 2 h. Strikingly, most of the SrtA mutant bacteria were killed within 10 min (Fig. 3A) at this concentration, whereas the killing of WT strain JRS4 was much slower. Thus, SrtA also contributes to the resistance against sPLA2-IIA by slowing down the killing rate of S. pyogenes.
To investigate whether this was also the case in our system, we compared the sensitivity of the JRS4 WT and SrtA
mutant bacteria to sPLA2-IIA in early stationary phase (OD$_{620}$ = 0.9) and mid-log phase (OD$_{620}$ = 0.4). As shown above, when analyzed in mid-log phase, the SrtA$^{-}$ mutant was sensitive to sPLA2-IIA; however, when it was analyzed in stationary phase, the SrtA$^{-}$ mutant was equally as resistant to sPLA2-IIA as JRS4 (Fig. 3B).

**FIGURE 2.** SrtA significantly contributes to the ability of *S. pyogenes* to resist the bactericidal activity of sPLA2-IIA. *A*, Overnight cultures of the WT strain JRS4 and the isogenic SrtA$^{-}$ mutant were diluted 20 times into fresh broth, and bacteria were grown to mid-log phase (OD$_{620}$ = 0.4). Bacteria were directly diluted 1000-fold into HEPES buffer to give a bacterial concentration of $\sim 1 \times 10^5$ CFU/ml. Recombinant human sPLA2-IIA was serially diluted in HEPES buffer, and small aliquots of 10 $\mu$l were added in triplicates to sterile 96-well polypropylene plates, after which it was mixed with 10 $\mu$l of the bacterial suspension. After incubation at 37°C in 5% CO$_2$ for 2 h, the samples were diluted in PBS, plated onto blood agar plates, and incubated overnight to determine the bacterial viability by colony counting. The killing is presented as the percentage of CFU for each bacterial strain remaining in the wells after incubation with sPLA2-IIA compared with the inoculum.

**FIGURE 2.** SrtA significantly contributes to the ability of *S. pyogenes* to resist the bactericidal activity of sPLA2-IIA. A, Overnight cultures of the WT strain JRS4 and the isogenic SrtA$^{-}$ mutant were diluted 20 times into fresh broth, and bacteria were grown to mid-log phase (OD$_{620}$ = 0.4). Bacteria were directly diluted 1000-fold into HEPES buffer to give a bacterial concentration of $\sim 1 \times 10^5$ CFU/ml. Recombinant human sPLA2-IIA was serially diluted in HEPES buffer, and small aliquots of 10 $\mu$l were added in triplicates to sterile 96-well polypropylene plates, after which it was mixed with 10 $\mu$l of the bacterial suspension. After incubation at 37°C in 5% CO$_2$ for 2 h, the samples were diluted in PBS, plated onto blood agar plates, and incubated overnight to determine the bacterial viability by colony counting. The killing is presented as the percentage of CFU for each bacterial strain remaining in the wells after incubation with sPLA2-IIA compared with the inoculum. B, Trans-complementation analysis where the SrtA$^{-}$ mutant strain was transformed with plasmid pJRS757 containing the intact srtA gene. The resulting trans-complemented strain (SrtA$^{-}$/pJRS757) was compared with the WT strain JRS4 and the SrtA$^{-}$ mutant in sensitivity to sPLA2-IIA using the protocol described above. C, The experiment was essentially performed as described in A, but the sPLA2-IIA inhibitor LY311727 was added to a final concentration of 50 $\mu$M.
This result showed that sPLA2-IIA is bactericidal only against growing bacteria and that SrtA most likely mediates protection of *S. pyogenes* against sPLA2-IIA at sites in the bacterial envelope engaged in growth.

**SrtA is necessary for S. pyogenes to resist sPLA2-IIA–mediated degradation of membrane phospholipids**

The bactericidal activity of sPLA2-IIA is due to its ability to degrade the bacterial membrane phospholipids. Thus, we examined the effect of sPLA2-IIA on [3H]-OA release from [3H]-OA–labeled JRS4 WT and SrtA– mutant bacteria in the bactericidal assay. As shown in Fig. 4, ∼14% of the total [3H]-OA was released from the SrtA– mutant, whereas only ∼2.5% of the total [3H]-OA was released from the WT JRS4 strain. Importantly, release of [3H]-OA from the SrtA– mutant was completely inhibited by LY311727, which inhibits the enzymatic activity of sPLA2-IIA, confirming that [3H]-OA release from the SrtA– mutant is due to degradation of phospholipids by sPLA2-IIA. This result showed that sPLA2-IIA degrades the membrane phospholipids to a greater extent in the SrtA– mutant compared with the parental strain JRS4, which clearly correlates with the bactericidal activity of sPLA2-IIA against the SrtA– mutant. Thus, SrtA protects *S. pyogenes* against degradation of the cellular membrane phospholipids by sPLA2-IIA.

**Human sPLA2-IIA shows greater binding to S. pyogenes WT bacteria compared with the SrtA– mutant**

Our data above showed that SrtA protects *S. pyogenes* from killing by sPLA2-IIA, indicating that one or more *S. pyogenes* LPXTG proteins are involved in the protection against this bactericidal molecule. We hypothesized that an LPXTG protein(s) contributes to the resistance to sPLA2-IIA by binding and sequestering sPLA2-IIA at the bacterial surface, thereby preventing it from reaching the cellular phospholipid membrane. Assuming that this hypothesis is correct, the WT JRS4 bacteria may show greater binding to sPLA2-IIA compared with the SrtA– mutant. To test this hypothesis, we analyzed the binding of radioiodinated sPLA2-IIA to JRS4 and the SrtA– mutant. As shown in Fig. 5A and 5B, sPLA2-IIA bound more avidly to the WT JRS4 bacteria compared with the SrtA– mutant. To exclude the possibility that the greater amount of sPLA2-IIA bound to JRS4 compared with the SrtA– mutant was due to differences in internalization of sPLA2-IIA rather than direct cell surface binding, we performed sPLA2-IIA–elution experiments with high ionic strength, a washing step that is known to release bound sPLA2-IIA from anionic components present at cellular surfaces (46). We incubated JRS4 with sPLA2-IIA, washed the bacteria, and subsequently eluted surface-bound sPLA2-IIA with 1 M NaCl. As shown in Fig. 5C, <10% of the bound sPLA2-IIA remained associated with JRS4 after incubation in 1 M NaCl, showing that almost all sPLA2-IIA was bound to the surface.

**sPLA2-IIA incubated with WT JRS4 bacteria retains its bactericidal properties**

*S. pyogenes* possess several LPXTG surface proteins that act as proteases, including the C5a peptidase and SpyCEP, which cleave and inactivate host defense molecules, such as complement C5a.
and IL-8 (42, 47, 48). To analyze whether sPLA₂-IIA is inactivated when bound to JRS4, we analyzed the bactericidal activity of sPLA₂-IIA against the GBS strain BM110 after elution from JRS4. As shown in Fig. 5D, sPLA₂-IIA eluted from JRS4 retained its bactericidal properties, because it was capable of killing GBS strain BM110 in a dose-dependent manner. Importantly, the killing was specifically mediated by sPLA₂-IIA, because it was capable of killing GBS strain BM110 in a dose-dependent manner. Importantly, the killing was completely inhibited by LY311727. Therefore, it is unlikely that the mechanism for protection against sPLA₂-IIA in JRS4 is by proteolytic cleavage and inhibition of the bactericidal activity of sPLA₂-IIA. This was further supported by the finding that the sPLA₂-IIA eluted from JRS4 did not appear to differ in size compared with that eluted from the SrtA mutant or the control sPLA₂-IIA not preincubated with bacteria (Fig. 5B).

SrtA does not protect S. pyogenes against positively charged AMPs

As mentioned above, D-alanylation of LTA was shown to contribute to resistance to both sPLA₂-IIA and positively charged AMPs. Because sPLA₂-IIA is a highly positively charged molecule, it is possible that the SrtA-mediated resistance mechanism against sPLA₂-IIA might also affect the sensitivity to AMPs, most of which are highly basic. To examine whether SrtA also contributes to the resistance to AMPs, we analyzed the difference in sensitivity between the WT JRS4 strain and the isogenic SrtA mutant to the positively charged human cathelicidin LL-37 and Magainin II. As shown in Fig. 6, there was little or no difference in sensitivity to LL-37 or Magainin II between JRS4 and the SrtA mutant, indicating that the SrtA-mediated resistance mechanism in S. pyogenes is specific for sPLA₂-IIA.

S. pyogenes SrtA is important for resistance against sPLA₂-IIA in vivo

To determine whether SrtA is important for S. pyogenes to be able to resist sPLA₂-IIA in vivo, we used Tg mice overexpressing human sPLA₂-IIA (Tg sPLA₂-IIA). These mice were generated in the WT C57BL/6 background, in which there is a natural mutation in the Pla2g2a gene encoding endogenous mouse sPLA₂-IIA (37). The Tg sPLA₂-IIA mice contain high levels of human sPLA₂-IIA activity in serum and various organs (Table I), whereas no sPLA₂-
Studies of microbial immune escape are of fundamental importance for understanding the mechanisms by which pathogens can establish and develop infections. Previous studies demonstrated two main mechanisms by which pathogenic bacteria resist or subvert sPLA₂-IIA. First, *B. anthracis* (17, 49) and *Bordetella pertussis* (5) were shown to express toxins that inhibit the secretion of sPLA₂-IIA from macrophages. Second, Gram-positive pathogens, such as *S. aureus*, evade sPLA₂-IIA attack by modifying the surface LTA with D-alanyl residues (18), which is believed to limit sPLA₂-IIA’s ability to cross the bacterial cell wall. In this article, we reported both in vitro and in vivo evidence that the transpeptidase SrtA increases *S. pyogenes*’ resistance against the bactericidal activity of human sPLA₂-IIA. Therefore, our results indicated that one or more LPXTG surface protein(s) in *S. pyogenes* are responsible for the protection against sPLA₂-IIA and represents a novel mechanism by which a Gram-positive bacterium can resist killing by sPLA₂-IIA.

In this study, we showed that clinical strains of *S. pyogenes* exhibit a high degree of resistance against sPLA₂-IIA. Compared with strains of GBS, which are sensitive to sPLA₂-IIA, an ~500-fold higher concentration was needed to kill 50% of the bacteria. The SrtAΔ mutation in *S. pyogenes* results in an ~30-fold increase in sensitivity against sPLA₂-IIA, showing that SrtA strongly affects the sensitivity to sPLA₂-IIA in *S. pyogenes*. However, SrtA cannot explain the entire difference in sensitivity compared with GBS, indicating that *S. pyogenes* also has other important means to evade sPLA₂-IIA compared with GBS. It is possible that *S. pyogenes* expresses other surface structures that contribute to the resistance.

Importantly, we demonstrated that SrtA contributes to the protection of *S. pyogenes* against sPLA₂-IIA but not positively charged AMPs, indicating that this mechanism is specific for protection against sPLA₂-IIA. We hypothesized that *S. pyogenes* use one or more LPXTG protein(s) that specifically bind and sequester sPLA₂-IIA at the bacterial surface, thereby preventing this molecule from reaching the cellular phospholipid membrane. Indeed, we demonstrated that WT *S. pyogenes* bacteria showed ~5-fold higher binding to sPLA₂-IIA compared with the SrtAΔ mutant, supporting this hypothesis. An alternative explanation for the greater association of sPLA₂-IIA to JRS4 compared with the SrtAΔ mutant could be that sPLA₂-IIA is internalized to a greater extent into the WT JRS4 bacteria. To exclude this possibility, we showed that >90% of the bound sPLA₂-IIA dissociated from the WT bacteria when incubated in a high-salt buffer, providing strong evidence that most sPLA₂-IIA is bound to the bacterial surface in JRS4. We also showed that sPLA₂-IIA bound to WT *S. pyogenes* retains its bactericidal properties, providing further evidence that the main mechanism to resist sPLA₂-IIA is to sequester this molecule at the bacterial surface.

To analyze the role of *S. pyogenes* SrtA in the resistance against sPLA₂-IIA–mediated killing in vivo, we developed a mouse model of mixed infection measuring the CI, which gives the relative contribution of individual bacterial factors to virulence (41, 50, 51). In this model, we showed that the ratio of JRS4/SrtAΔ bacteria found in liver and spleen is significantly higher in the output from Tg sPLA₂-IIA mice compared with WT mice lacking expression of sPLA₂-IIA, most likely by protecting the bacteria against direct killing by sPLA₂-IIA. Importantly, because there was no apparent difference in inflammatory status between Tg sPLA₂-IIA and WT C57BL/6 mice.
mice during the experiment, we can rule out that the difference in CI is due to different levels of inflammation between the two types of mice. Of note, the concentration of human sPLA2-IIA in serum from Tg sPLA2-IIA mice (∼2500 μg/l) is comparable to levels of sPLA2-IIA measured in serum from human patients with severe sepsis (880 μg/l) or typhoid fever (1440 μg/l) (14, 52, 53), indicating that the level of human sPLA2-IIA in Tg mice is physiologically relevant.

Other investigators have successfully used SrtA mut mutants as the basis for analysis of the role of individual LPXTG proteins in interactions between Gram-positive pathogens and their infected host. For example, in a mouse model of oral infection, Lalioui et al. (54) showed that an SrtA mut strain of GBS was strongly impaired in ability to colonize the intestine, a location that is a natural habitat for GBS in humans. In a subsequent study, it was demonstrated that a single LPXTG protein, HvgA, is responsible for the ability of GBS to colonize the gut (55). In another study, Cheng et al. (56) used an SrtA mut mutant of S. aureus in analyses demonstrating that LPXTG proteins were required for this pathogen to cause kidney abscess formation in the mouse model, which led to the identification of the LPXTG protein SdrD as the main virulence factor in this model. Thus, the use of SrtA mut mutants as the basis for studies on the interaction between Gram-positive pathogens and their host has revealed important functions for individual LPXTG proteins. Among the LPXTG proteins in S. pyogenes are several potential virulence factors that could be responsible for the resistance to sPLA2-IIA. Although we cannot rule out that more than one LPXTG protein contributes to the resistance, we hypothesize that a single LPXTG protein is responsible for the protection against sPLA2-IIA. Indeed, the ligand-binding properties of LPXTG proteins in interactions with host proteins very often are highly specific, with well-defined binding domains in the bacterial proteins (26, 40, 42, 57, 58). Although S. pyogenes is a clinically important pathogen, we still know little about the in vivo role of different antibacterial molecules in the defense against S. pyogenes. In a study by Nizet et al. (59), it was shown that mice deficient in the cathelicidin CRAMP, the mouse equivalent of LL-37, are more sensitive to skin infections caused by S. pyogenes. A recent study by Shannon et al. (60) showed that the histidine-rich glycoprotein, an abundant plasma protein, protects mice against S. pyogenes infection by the s.c. route, a property that may be due, at least in part, to its antibacterial activity. In this study, we showed that Tg mice overexpressing human sPLA2-IIA are significantly protected against lethal S. pyogenes infection compared with WT C57BL/6 mice. This result showed that human sPLA2-IIA contributes to the host defense against serious experimental S. pyogenes infection and provides a novel example of an antibacterial molecule that may be involved in the protection against S. pyogenes infections in humans.
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Disclosures
The authors have no financial conflicts of interest.

References


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