1 SUPPLEMENTAL METHODS

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3 Immunogenicity assessment. To assess the humoral immune response to SAL200, the serum 4 level of antibodies against rSAL-1 was determined using a conventional bridging ELISA with streptavidin-biotin detection. As an antibody standard, anti-rSAL-1 immunoglobulins (2 5 6 mg/ml) were prepared according to the conventional IgG purification method from immune 7 sera obtained from dogs immunized with SAL200. The prepared anti-rSAL-1 8 immunoglobulins were confirmed via immunodepletion testing. Biotinylated rSAL-1 (1 mg/ml) was obtained from Younginfrontier (Seoul, Republic of Korea). The streptavidin-HRP 9 solution (1 mg/ml) was obtained from Sigma-Aldrich (St. Louis, MO, USA). An ELISA 10 11 starter kit was obtained from Koma Biotech (Seoul, Republic of Korea). ELISA was performed using a conventional method with modification. Briefly, SAL200 was diluted to 1 12 mg/ml in a formulation buffer (1.56 g/l L-histidine [pH 6.0], 50 g/l D-sorbitol, 1.47 g/l 13 CaCl₂·2H₂O, and 1 g/l poloxamer 188). The diluted SAL200 solution was further diluted to 1 14 µg/ml using the coating buffer (0.05 M sodium carbonate buffer, pH 9.6) provided in the 15 16 ELISA starter kit. The prepared solution (100 µl) was added to each well of an ELISA plate and then covered with a plate sealer, after which the plate was incubated at 37°C for 1 h. The 17 plate was subsequently washed three times by applying 250 µl of washing buffer (phosphate-18 buffered saline [PBS] containing 0.05% Tween-20) per well. For blocking, 200 µl of 19 blocking buffer (PBS containing 0.05% Tween-20, 0.5% bovine serum albumin, and 20% 20 fetal bovine serum) was added to each well to block the remaining binding sites on the plate. 21 22 After the addition of the blocking buffer, the plate was incubated for 2 h at 37°C. Next, the wells were washed as described above. ELISA samples, including calibration standards, 23 quality control standards, and test samples, were prepared during the blocking and washing 24

25 steps. Calibration standards, including anchor points (0.781, 1.563, 3.125, 6.25, 12.5, 25, 50, and 100 µg/ml) and quality control standards (2, 10, and 40 µg/ml), were prepared using the 26 anti-rSAL-1 antibody standard and human pooled serum. The calibration standards, quality 27 control standards, and test samples were diluted 1:20 in PBS, and the mixtures were briefly 28 vortexed. The prepared ELISA samples (100 µl) were added to the wells of the plate and 29 shaken for 1 min using an orbital shaker. The plate was subsequently incubated for 1 h at 30 23°C. After incubation, the wells were washed again as described above. A solution of 31 32 1:10,000 biotinylated rSAL-1 was prepared in a dilution buffer (PBS containing 0.05% Tween-20, 0.5% bovine serum albumin, and 10% fetal bovine serum). Then, 100 µl of the 33 diluted biotinylated rSAL-1 solution was added to each well of the plate, and the plate was 34 35 covered with a plate sealer. The plate was then incubated for 1 h at 23°C, and the wells were re-washed. A 1:10,000 streptavidin-HRP solution was prepared in a dilution buffer (PBS 36 containing 0.05% Tween-20 and 0.5% bovine serum albumin). The diluted streptavidin-HRP 37 solution (100 μ l) was added to each well, and the plate was covered with a plate sealer. The 38 plate was incubated for 1 h at 23°C. After incubation, the wells were re-washed. Next, 100 µl 39 40 of the provided tetramethylbenzidine (TMB) substrate solution was added to each well, and 41 the plate was incubated for no longer than 30 min at room temperature in the dark. The reaction was stopped by the addition of 100 μ l of stop solution, and the OD₄₅₀ was recorded. 42 43 The ELISA was effective in an analysis range of 1.563-50 µg/ml, and the within-run accuracy, within-run precision, between-run accuracy, and between-run precision values were 44 95.49-107.5%, 1.770-4.919%, 94.50-110.0%, and 2.342-7.012%, respectively. 45

46 Determination of rSAL-1 levels in serum. For the pharmacokinetic analyses, the
47 rSAL-1 levels in serum were determined using a validated double antibody sandwich ELISA.
48 A monoclonal anti-rSAL-1 antibody and a monoclonal anti-rSAL-1 antibody conjugated to

49 HRP were obtained from Younginfrontier (Seoul, Republic of Korea). An ELISA starter kit was obtained from Koma Biotech (Seoul, Republic of Korea). ELISA was performed 50 according to a conventional method with modification. Briefly, 1:400 dilutions of a 2-mg/ml 51 solution of monoclonal anti-rSAL-1 antibody (capture antibody) were prepared with a 52 coating buffer (0.05 M sodium carbonate buffer, pH 9.6). The diluted capture antibody 53 54 solution (100 µl) was added to each well of the ELISA plate and then covered with a plate sealer, after which the plate was incubated overnight at 4°C. The plate was subsequently 55 56 washed five times by applying 250 µl of a washing buffer (Tris-buffered saline [TBS] containing 0.05% Tween-20) per well. For blocking, 200 µl of blocking buffer I (Applichem, 57 Darmstadt, Germany) was added to each well to block the remaining binding sites on the 58 59 plate. After the addition of the blocking buffer, the plate was incubated for 1 h at 37°C. Next, the wells were washed as described above. ELISA samples, including calibration standards, 60 quality control standards, and test samples, were prepared during the blocking and washing 61 steps. Calibration standards, including anchor point (0.008, 0.016, 0.031, 0.063, 0.125, 0.250, 62 and 0.500 µg/ml) and quality control standards (0.048, 0.095, and 0.350 µg/ml), were 63 64 prepared using SAL200 and human pooled serum. The calibration standards, quality control standards, and test samples were then diluted 1:10 in TBS containing 5.5 mM CaCl₂, and the 65 mixtures were briefly vortexed. The prepared ELISA samples (100 µl) were added to the 66 67 wells of the plate and shaken for 1 min using an orbital shaker. The plate was subsequently incubated for 3 h at 25°C. After incubation, the wells were re-washed as described above. A 68 2.5-mg/ml solution of monoclonal anti-rSAL-1 antibody conjugated to HRP (detection 69 70 antibody) was diluted 1:5,000 in TBS. After re-washing the plate, 100 µl of the diluted detection antibody solution was added to each well of the plate, and the plate was covered 71 with a plate sealer. The plate was then incubated for 2 h at 25°C, and the wells were re-72

washed. Next, 100 μ l of the TMB substrate solution was added to each well, and the plate was incubated for no longer than 30 min at room temperature in the dark. The reaction was stopped by the addition of 100 μ l of stop solution, and the OD₄₅₀ was recorded. This ELISA method was effective in an analysis range of 0.016-0.500 μ g/ml, and the within-run accuracy, within-run precision, between-run accuracy, and between-run precision values were 96.88-116.7%, 2.259-15.67%, 93.92-103.4%, and 8.670-12.15%, respectively.

79 **Pharmacokinetic analysis.** Pharmacokinetic calculations were performed using the pharmacokinetic program PhoenixTM WinNonlin[®] version 6.3 (Pharsight, Mountain View, 80 CA, USA) with non-compartmental methods. The C_{max} and t_{max} values were obtained directly 81 from the serum concentration-time profiles. The $t_{1/2}$ value over the sampling period was 82 83 calculated as the mean residence time from the time of dosing to the time of the last quantifiable concentration (MRT_{last}): MRT_{last} \times 0.693. MRT_{last} was calculated as AUMC_{last} / 84 85 AUC_{last}, where AUMC_{last} denotes the area under the first moment curve from time zero to the time at which the last sample with a measurable concentration was obtained. The area under 86 the serum concentration-time curve from time zero to the time at which the last sample with a 87 measurable concentration (AUC_{last}) was calculated using the linear trapezoidal rule for the 88 89 ascending region of the concentration-time curve and the log-linear trapezoidal rule for the descending region of the concentration-time curve. Data below the LLOQ were excluded 90 91 from the AUC_{last} calculation. The AUC from time zero to infinity (AUC_{inf}) was estimated as 92 $AUC_{last} + C_{last} / K_{el}$, where C_{last} denotes the concentration in the last serum sample at which a measurable concentration was obtained. The total body clearance (CL) was calculated as the 93 94 dose / AUC_{inf}. All data were expressed as the means \pm SDs.

95 Pharmacodynamic assay. Pharmacodynamics were measured using an *ex vivo*96 blood assay for determining blood antibacterial activity. Serum samples collected after

97 intravenous dosing were used in the ex vivo blood assay. Briefly, 20 µl of S. aureus glycerol stock was inoculated into a 50-ml conical tube containing 20 ml of tryptic soy broth (TSB; 17 98 g/l casein digest, 3 g/l soybean digest, 2.5 g/l dextrose, 5 g/l NaCl, and 2.5 g/l dipotassium 99 phosphate) and incubated overnight at 37°C. Next, 20 µl of the overnight bacterial culture 100 101 was added to a 50-ml conical tube containing 20 ml of TSB medium. The conical tube was incubated at 37°C until the OD₆₀₀ reached 1.8-2.0, at which time 2 ml of culture was 102 distributed evenly across the surface of a tryptic soy agar (TSA; 15 g/l casein digest, 5 g/l 103 104 soybean digest, 5 g/l NaCl, and 15 g/l agar) plate, and the excess broth was removed after standing for 5 min. The inoculated plate was then allowed to dry in a laminar flow hood for 105 approximately 30 min with the lid ajar. After drying, the plate was incubated overnight at 106 107 37° C. Standards (0, 0.05, 0.1, 0.2, 0.4, 0.6, 0.8, and 1 µg/ml) were subsequently prepared by spiking SAL200 into human pooled serum containing 10 mM CaCl₂. Test samples were 108 prepared by adding a 1% (v/v) volume of 1 M CaCl₂ to the serum samples collected from 109 each participant. Then, 10-µl aliquots of standard and test samples were spotted onto the plate. 110 The plate was allowed to dry in a laminar flow hood for approximately 10 min with the lid 111 The plate was incubated overnight at 37°C. Following incubation, the 112 ajar. pharmacodynamics were accessed via a qualitative comparison of the clear zones (i.e., lysis 113 halos) formed after spotting the test samples onto the lawn of S. aureus with those formed 114 115 after spotting the standards.

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121 SUPPLEMENTAL TABLE

122 Table S1. Adverse events after a single intravenous dose among healthy male volunteers

123 (cohort 5, individual)

Participant			Observed time				Comment
identification		Observed adverse					(number of
code (dose,	Dosing	event (number of	Day	Day	Day	\geq Day	adverse
mg/kg; A/P ^a)	date	adverse events)	1^{b}	2	3	7	events)
R501 (10; A)	Nov 12,	Rigors (1)	•				Related (1)
	2013	Dizziness (1)	•				Related (1)
		Myalgia (2)	٠			٠	Related (2)
		Headache (1)	٠				Related (1)
		Abdominal pain (1)		•			Related (1)
		Fatigue (1)		•			Related (1)
		Pharyngitis (1)			•		Related (1)
		Rhinitis (1)				•	Related (1)
R502 (10; A)	Nov 12,	Nausea (1)	٠				Related (1)
	2013	Dizziness (1)	•				Related (1)
		Rigors (1)	٠				Related (1)
		Fatigue (1)	•				Related (1)
		Myalgia (1)	•				Related (1)
		Hypotension (1)	•				Related (1)
R503 (10; A)	Nov 12,	Rigors (1)	•				Related (1)
	2013	Headache (1)	•				Related (1)
		Back pain (1)	•				Related (1)

		Nausea (1)	•		Related (1)
		Fatigue (1)	•		Related (1)
		Dyspepsia (1)	•		Related (1)
R504 (10; P)	Nov 19,	None			
	2013				
R505 (10; A)	Nov 19,	Rigors (1)	•		Related (1)
	2013	Headache (1)	•		Related (1)
R506 (10; A)	Nov 26,	Rhinitis (2)	• •		Related (1)/
	2013				Unrelated (1)
R507 (10; P)	Nov 26,	Headache (3)	•	•/•	Related (2)/
	2013				Unrelated (1)
R508 (10; A)	Nov 26,	Fever (1)	•		Related (1)
	2013	Rhinitis (1)		•	Unrelated (1)
		Coughing (1)		•	Unrelated (1)

124 ^aA, active; P, placebo

^bDosing day