

33 **Abstract:**

34 The seriousness of microbial resistance combined with the lack of new antimicrobials have
35 increased the interest in the development of antimicrobial peptides (AMPs) as novel therapeutics.
36 In this study, we evaluated the antimicrobial activity of two short synthetic peptides, namely
37 RRIKA and RR. These peptides exhibited potent antimicrobial activity against *Staphylococcus*
38 *aureus* and their antimicrobial effects were significantly enhanced by addition of three amino
39 acids in the C terminus, which consequently increased the amphipathicity, hydrophobicity and
40 net charge. Moreover, RRIKA and RR demonstrated a significant and rapid bactericidal effect
41 against clinical and drug-resistant *Staphylococcus* isolates including methicillin-resistant
42 *Staphylococcus aureus* (MRSA), vancomycin-intermediate *S. aureus* (VISA), vancomycin-
43 resistant *S. aureus* (VRSA), linezolid-resistant *S. aureus* and methicillin-resistant *S. epidermidis*.
44 In contrast to many natural AMPs; RRIKA and RR retained their activity in the presence of
45 physiological concentrations of NaCl and MgCl₂. Both RRIKA and RR enhanced the killing of
46 lysostaphin over 1000-fold and eradicated MRSA and VRSA isolates within 20 minutes.
47 Furthermore, the peptides presented were superior in reducing adherent biofilms of *S. aureus* and
48 *S. epidermidis* when compared to conventional antibiotics. Our findings indicate that the
49 staphylocidal effects of our peptides were through permeabilization of the bacterial membrane,
50 leading to leakage of cytoplasmic contents and cell death. Furthermore, peptides were not toxic
51 to HeLa cells at 4 to 8 fold their antimicrobial concentrations. The potent and salt-insensitive
52 antimicrobial activities of these peptides present an attractive therapeutic candidate for treatment
53 of multidrug-resistant *S. aureus* infections.

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55

56 **INTRODUCTION**

57 Methicillin-resistant *Staphylococcus aureus* (MRSA) infections have become a major public
58 health concern in both hospital and community settings (1, 2). Since the late 1990s, community-
59 associated MRSA (CA-MRSA) emerged as a principal cause of skin and soft-tissue epidemics
60 throughout the world (3). It has been estimated that mortality rates from MRSA infections in US
61 hospitals are higher than infections caused by HIV/AIDS and tuberculosis combined (4).
62 Furthermore, there is a considerable medical challenge for treating MRSA infections due to the
63 remarkable ability of *S. aureus* to develop resistance to multiple antibiotics, thus limiting the
64 number of viable therapeutic options (2, 5). Therefore, there is an urgent need to develop novel
65 antimicrobials with unique mechanisms of action to combat MRSA.

66 Antimicrobial peptides (AMPs) are key components of the host innate defense against infections
67 in most creatures (6, 7). The potential therapeutic applications of AMPs are significant (8). They
68 have rapid and broad spectrum antibacterial activity. In addition, it is extremely difficult for
69 bacteria to evolve resistance against AMPs that disrupt the microbial membrane, as it would
70 necessitate fundamental alterations in the lipid composition of the bacterial membranes. Though
71 AMPs have many important antibacterial properties, several studies have exposed their potential
72 limitations as therapeutic agents.

73 A significant number of natural AMPs are large in size with high host cytotoxicity and moderate
74 antimicrobial activity. Moreover, their production cost is high (8, 9). In addition, many natural
75 AMPs lose their antimicrobial activity in physiological salt concentrations (10, 11). These
76 characteristics have substantially hindered their pharmaceutical development as new therapeutic
77 agents. Thus, successful development of novel AMPs as future therapeutics requires

78 identification of short AMPs demonstrating strong antimicrobial activity, salt tolerance and
79 minimal toxicity to host tissues.

80 Previously, our group has developed short synthetic peptide templates for therapeutic
81 applications against hyperplasia and inflammation (12-14). However their antimicrobial activity
82 was never explored. In the present study, we screened four synthetic peptides and identified two
83 peptides, namely RRIKA and RR, with potent antimicrobial activity against a panel of clinical
84 and drug resistant *Staphylococci* strains. In addition, we investigated the synergistic activity of
85 these peptides in combination with clinically relevant antimicrobials and examined the peptides'
86 ability to disrupt staphylococcal biofilms. Moreover, we performed a series of experiments to
87 explore the antibacterial mechanism of action of RRIKA and RR and examined the toxicity of
88 these peptides toward mammalian cells.

89 **MATERIALS AND METHODS**

90 **Bacterial strains and media**

91 All *Staphylococcus* strains tested in this study are presented in table (1). Mueller-Hinton broth
92 (MHB) and Mueller-Hinton agar (MHA) were purchased from Sigma-Aldrich, while trypticase
93 soy broth (TSB) and trypticase soy agar (TSA) were purchased from Becton-Dickinson,
94 Cockeysville, Md.

95 **Reagents, peptides and antibiotics**

96 Nisin (Sigma, N5764), melittin from honey bee venom (Sigma, M2272), magainin I (Sigma,
97 M7152), vancomycin hydrochloride (Gold Biotechnology), linezolid (Selleck Chemicals),
98 ampicillin sodium salt (IBI Scientific), recombinant lysostaphin (3000 U/ mg)

99 from *Staphylococcus simulans* (Sigma, L9043) and calcein AM (Molecular Probes, Life
100 Technologies) were all purchased from commercial vendors.

101 **Peptide synthesis**

102 Peptides RRIKA, RR, KAF, and FAK were synthesized on Knorr-amide resin (Synbiosci Corp.)
103 using standard Fmoc chemistry. Two different chemistries were used to couple each amino
104 acid. The first coupling reagents were N-hydroxybenzotriazole (HOBt)/N, N'-
105 diisopropylcarbodiimide (DIC) and the second coupling reagents were 2-(1Hbenzotriazole-1-yl)-
106 1, 1,3,3-tetramethyluronium hexafluorophosphate (HBTU) and lutidine. For FITC labeled
107 YARA, an aminohexanoic acid spacer was added to the N-terminus to serve as a spacer for the
108 addition of FITC isomer 1 (Molecular Probes). The FITC isomer was solubilized in 12:7:5
109 pyridine/DMF/DCM and incubated with the deprotected peptide overnight. A ninhydrin test was
110 used to check complete coupling of FITC to the peptide. Following synthesis, the peptide was
111 cleaved from the resin with a trifluoroacetic acid-based cocktail, precipitated in ether, and
112 recovered by centrifugation. The recovered peptide was dried in vacuo, resuspended in MilliQ
113 purified water, and purified using an FPLC (ÅKTA Explorer, GE Healthcare) equipped with a
114 22/250 C18 prep-scale column (Grace Davidson). An acetonitrile gradient with a constant
115 concentration of 0.1% trifluoroacetic acid was used to achieve purification. Desired molecular
116 weight was confirmed by time-of-flight MALDI mass spectrometry using a 4800 Plus MALDI
117 TOF/TOF™ Analyzer (Applied Biosystems).

118 **Antibacterial assays**

119 Minimum inhibitory concentrations (MIC) of peptides and antibiotics were evaluated with the
120 broth microdilution technique in MHB with an initial inoculum of 5×10^5 cells non-treated
121 Polystyrene microtiter plates (CytoOne, CC7672-7596) in accordance with the Clinical and

122 Laboratory Standards Institute (CLSI) (15). The MIC was interpreted as the lowest concentration
123 of peptide or antibiotic that completely inhibited the visible growth of bacteria after 16 hr
124 incubation of the plates at 37°C. Each agent was tested in triplicate in at least two independent
125 experiments. The highest MIC value was reported.

126 **Antimicrobial activity in the presence of salts**

127 To investigate the activity of peptides in the presence of high salt concentrations, the MIC was
128 again identified as described above except, fixed concentrations of NaCl or MgCl₂ were added to
129 each well of the microtiter plate. For further analysis of the effect of salinity on killing ability of
130 peptides; logarithmic phase of MRSA USA300 and VRSA VRS13 at 5×10^5 colony forming
131 units/ml (CFU/ml) were incubated with $2 \times$ MIC of RRIKA and RR, in MHB, with different
132 concentrations of NaCl (0, 50, 100, 150 mM) or MgCl₂ (0, 1, 2, 3 mM) for 4 hr. Bacteria were
133 serially diluted and plated, in triplicate, on TSA plates. CFUs were counted after incubation of
134 plates for 24 hr at 37°C.

135 **Time-kill assay**

136 An overnight culture of MRSA USA300 was diluted in fresh MHB and incubated until optical
137 density (OD₆₀₀) ≈ 1 , was reached, then washed twice with phosphate buffered saline (PBS)
138 diluted to $\sim 5 \times 10^5$ CFU/ml in MHB. Peptides, vancomycin and linezolid were added at
139 concentrations equivalent to $5 \times$ MIC. Peptide diluent (sterile water) served as a negative control.
140 Bacterial cell viability was monitored up to 24 hr. Samples were removed at specific intervals,
141 serially diluted in PBS, and plated, in triplicate, on TSA plates. CFUs were counted after 24 hr
142 incubation of plates at 37°C. To study the effect of serum and salinity on the killing kinetics of
143 the peptides, a time-kill assay was performed as described above with the exception that bacteria

144 and peptides at $5 \times$ MIC were incubated in MHB alone or with 10% fetal bovine serum (FBS) or
145 incubated in PBS (containing 137 mM NaCl).

146 **Synergy with antimicrobials**

147 The fractional inhibitory concentration (FIC) index was utilized to determine the relationship
148 between antimicrobial agents. Peptide MICs against test microorganisms were determined in
149 triplicate samples. Two-fold serial dilutions of antimicrobials (lysostaphin, vancomycin,
150 linezolid and nisin) were tested in the presence of a fixed concentration of peptide, equals to $\frac{1}{4} \times$
151 peptide MIC. It is worthy to note that none of the peptides killed the test microorganisms at their
152 quarter MICs. The FIC index was calculated as follows: $FIC = 0.25 + MIC$ (antibiotic in
153 combination)/MIC (antibiotic alone); 0.25 is equal to MIC (peptide in combination)/MIC
154 (peptide alone). An FIC index of ≤ 0.5 is considered to demonstrate synergy. Additive effect was
155 defined as an FIC index of 1. Antagonism was defined as an FIC index of > 4 (16).

156 To further investigate the synergism of peptides with lysostaphin, the time-kill method of
157 determining synergy was utilized as described elsewhere (17). MRSA USA300 and VRSA
158 VRS13 were incubated with $\frac{1}{2} \times$ MIC of RRIKA or $\frac{1}{2} \times$ MIC of RR alone; or in combination
159 with $\frac{1}{4} \times$ MIC of lysostaphin (MIC of lysostaphin against MRSA USA300 and VRSA VRS13
160 were 0.04 and 0.08 μ M, respectively). Samples were obtained at different time points, then
161 diluted and plated on TSA plates. The plates were incubated for 24 hr at 37°C before CFU were
162 determined. Synergy was defined as a 100-fold or 2-log_{10} decrease in colony count by the
163 combination of two agents together as compared to each agent tested alone. Additivity or
164 indifference was defined as a 10-fold change in CFU and antagonism was defined as a 100-fold
165 increase in CFU by the combined peptide treatment in comparison to the single treatment.

166

167 **Bacterial membrane disruption activity (Bacteriolysis)**

168 Cell lysis, as indicated by a decrease in OD₅₉₅, was determined as described before (18). Briefly,
169 MHB was inoculated with an overnight culture of MRSA USA300 and incubated aerobically at
170 37°C until an OD₅₉₅≈0.6 was reached and then diluted to OD₅₉₅≈0.2 in MHB (equivalent to ≈10⁸
171 CFU/mL). 200 μL of diluted bacteria was added to all wells of a 96-well plate. RRIKA and RR
172 were added in concentrations equivalent to 4 × MIC. Nisin at 4 × MIC was used as a positive
173 control and untreated bacteria served as a negative control. MHB with the same concentrations
174 of drugs served as blanks. Turbidity was monitored at defined intervals up to 10 hr in a
175 Molecular Devices Vmax microplate reader at 595 nm absorption. The assay was carried out in
176 triplicate samples for each treatment regimen.

177 **Calcein leakage assay**

178 Membrane permeabilization of *S. aureus* by peptides was monitored and quantified by the
179 leakage of the preloaded fluorescent dye, calcein, as described before (19). MRSA USA300 was
180 grown in MHB to logarithmic phase at 37°C. Cells were then harvested by centrifugation,
181 washed twice with PBS, and then adjusted spectrophotometrically to an OD₆₀₀ of 1.0 (≈10⁹
182 CFU/ml) in PBS containing 10% (vol/vol) MHB. Then MRSA cells were incubated with 3 μM
183 calcein AM for 1 hr at 37°C. Calcein-loaded cells were harvested by centrifugation (3,000 × g,
184 10 min), suspended in PBS, and diluted to achieve a final inoculum of 10⁷ CFU/ml. Aliquots of
185 100 μL were then added into a sterile black-wall 96-well plate. RRIKA and RR were added in
186 concentrations equivalent to 5 × and 10 × MIC. Bacteria treated with peptide diluent (sterile
187 water) and nisin (10 μg/mL) served as negative and positive controls, respectively. Calcein
188 leakage was measured for 120 min using a fluorescence plate reader (FLx800 model BioTek®
189 Instruments, Inc. Winooski, Vermont). Membrane permeabilization (%) was calculated as the

190 absolute percent calcein leakage by peptides with respect to calcein-loaded with no-peptide
191 treated cells. Experiments were done in triplicate and repeated independently twice.

192 **DNA binding assay**

193 The ability of RRIKA and RR to bind DNA was investigated by the electrophoretic mobility
194 shift assay as has been previously described (20). In brief, increasing concentrations of RRIKA,
195 RR and magainin were incubated with 250 ng of plasmid DNA (pUC19) in 30 μ L binding buffer
196 (10 mM Tris-HCl, 1 mM EDTA buffer, pH 8.0) at room temperature for 30 min. After
197 incubation, the DNA was analyzed by 1% gel electrophoresis. DNA migration was visualized by
198 ethidium bromide staining.

199 **Quantification of activity against biofilms**

200 Biofilm-forming clinical isolates of *S. aureus* (ATCC 6538) and *S. epidermidis* (ATCC 35984)
201 were grown overnight in TSB. After incubation, cultures were diluted 1:100 in TSB
202 supplemented with 1% glucose. Diluted bacteria were inoculated in either 24- or 96-well flat
203 bottom cell culture plates (polystyrene) and incubated at 37°C for 24 hr. The culture medium was
204 subsequently removed and wells were carefully washed with PBS three times to remove
205 planktonic bacteria before re-filling wells with fresh MHB. Peptides and antibiotics were added
206 at different concentrations and plates were incubated at 37°C for 24 hr. After the removal of
207 medium at the end of incubation, wells were rinsed by submerging the entire plate in a tub
208 containing tap water. Biofilms were stained with 0.1% (w/v) crystal violet for 30 min. After
209 staining, the dye was removed and the wells washed with water. The plates were dried for at
210 least 1 hr, prior to addition of ethanol (95%) to solubilize the dye bound to the biofilm. The
211 optical densities (OD) of biofilms were measured at 595 nm absorbance by using a microplate

212 reader (Bio-Tek Instruments Inc.). An inverted microscope (Vista Vision, VWR), with attached
213 camera and 25X objective was used to photograph the biofilm in 24-well plates.

214 **Hemolysis assay**

215 Human red blood cells (RBCs) (Innovative Research “Novi, MI”) were pelleted by
216 centrifugation at 2000 rpm for 5 min followed by washing three times with PBS. An 8%
217 suspension (v/v) of RBCs was prepared in PBS and 50 μ L of the solution was transferred to a
218 96-well plate. Then, 50 μ L of different concentration of peptides in PBS were added to give a
219 final suspension of 4% (v/v) of RBCs. PBS served as a negative control. 0.1 % triton X-100 as
220 well as 5 μ M melittin served as a positive control. The plate was incubated at 37°C for 1 hr.
221 The plate was subsequently centrifuged at 1000 rpm for 5 min at 4°C. 75 μ L aliquots of the
222 supernatants in each well were carefully transferred to a new sterile 96-well plate. Finally the
223 hemolytic activity was evaluated by measuring the optical absorbance at OD₄₀₅ with a microplate
224 reader (Vmax, Molecular Devices). The hemolysis percentage was calculated based on the 100%
225 release with 0.1 % triton X-100 or 5 μ M melittin. Experiments were done in triplicates.

226 **Cytotoxicity assay**

227 The toxicity of RRIKA, RR and melittin against HeLa cells was carried out using the CellTiter
228 96[®] AQueous Non-Radioactive Cell Proliferation Assay kit from Promega (21). Briefly, $\sim 2 \times 10^4$
229 HeLa cells suspended in 200 μ L of DMEM supplemented with 10% FBS, were seeded in 96-
230 well plates and incubated at 37°C in a 5% CO₂ atmosphere for 24 hr. The HeLa cells were further
231 incubated with different concentration of peptides for 24 hr. At the end of the incubation period,
232 the culture media was discarded and the cells in each well were washed with PBS. 100 μ L of cell
233 culture media and 20 μ L CellTiter 96[®] AQueous assay reagent were added next (solution reagent

234 contains a tetrazolium compound [3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-
235 (4-sulfophenyl)-2H-tetrazolium, inner salt; MTS]). The plate was returned to the incubator for 2
236 hr to allow color development. The intensity of color was quantified at 490 nm using a 96-well
237 microplate reader (Vmax, Molecular Devices). Results were expressed as the percentage mean
238 absorbance by cells upon incubation with peptide with respect to incubation with the control
239 (untreated wells).

240 **Statistical analyses**

241 The mean absorbance of crystal violet in the biofilm reduction was compared between the groups
242 using the two-tailed Student *t* test ($p \leq 0.05$). All statistical analyses were performed with
243 Microsoft Excel's statistical program.

244 **RESULTS:**

245 **Antimicrobial activity**

246 The amino acid sequences and characteristics of the synthetic peptides used in this study are
247 shown in Table 2. We screened the synthetic peptides against a panel of clinical and drug-
248 resistant *Staphylococcal* strains (Table 1). RRIKA inhibited the growth of methicillin-sensitive
249 *S. aureus* (MSSA), MRSA, VISA, VRSA, linezolid-resistant *S. aureus*, methicillin-resistant *S.*
250 *epidermidis* and multidrug-resistant strains from concentrations ranging from 2 to 4 μ M. RR
251 inhibited the same isolates with MICs ranging from 8 to 32 μ M. In contrast, KAF and FAK
252 peptides were inactive against all strains tested up to 64 μ M (data not shown).

253 **Bacterial killing kinetics**

254 Figure 1 presents the rate of microbial killing by RR, RRIKA, vancomycin and linezolid when
255 MRSA USA300 was exposed to $5 \times$ MIC of each treatment over a 24 hr incubation period at 37

256 °C. Both RRIKA and RR exhibited a rapid bactericidal effect with 3- \log_{10} reduction (99.9 %
257 clearance) within 60 and 90 min, respectively. In addition, both peptides completely eliminated
258 the inoculum (5×10^5 CFU/mL) within 120 and 180 min, respectively. In comparison,
259 vancomycin achieved a 3- \log_{10} bacterial reduction only after 24 hr; while linezolid (a protein
260 synthesis inhibitor) exhibited a bacteriostatic effect producing only a single log bacterial
261 reduction over the 24 hr incubation. The killing kinetics of RRIKA and RR in PBS (containing
262 137mM NaCl) did not significantly change when compared to MHB (figure 2). In the presence
263 of 10% FBS, RRIKA retained its bactericidal activity, though it required longer to kill the
264 bacteria completely (6h), while RR lost its antibacterial activity at the same condition.

265 **Antimicrobial activity in the presence of salts**

266 The MIC of RRIKA and RR for MRSA USA300 was determined in the presence of NaCl or
267 MgCl_2 . There was no increase in the MICs of RRIKA and only a one fold increase in case of RR
268 was observed in the presence of 100 mM NaCl. Moreover, the antimicrobial activity of RRIKA
269 and RR was not impeded in the presence of MgCl_2 at a concentration equivalent to reported
270 physiological conditions (2 mM). To study the effect of salinity on the killing ability of the
271 peptides, CFU were counted after incubation of MRSA USA300 and VRSA VRS13 strains with
272 RRIKA and RR ($2 \times \text{MIC}$) in increasing concentrations of NaCl or MgCl_2 . As shown in
273 supplementary figure 1, there was no significant difference in the bactericidal activity of RRIKA
274 in the presence of NaCl up to 150 mM. Similar patterns were observed for RR with the exception
275 of one \log_{10} increase in CFU observed at 150 mM NaCl; however this peptide retained its
276 bactericidal activity at this concentration (producing a 3- \log_{10} reduction). Furthermore,
277 physiological concentrations of MgCl_2 (1, 2 and 3 mM) had no effect on the killing ability of
278 both RRIKA and RR.

279 **Synergy with antimicrobial agents**

280 Both RRIKA and RR acted synergistically with lysostaphin against MSSA, MRSA and VRSA
281 strains with a FIC index equal to 0.26 (Table 3). When both peptides were combined together,
282 only one fold decrease in each peptide's MIC was observed (FIC index = 0.75). However, an
283 additive effect was observed when each peptide was combined with vancomycin, linezolid or
284 nisin against tested strains (data not shown). There were no antagonistic effects observed
285 between the peptides and antimicrobials against all strains tested.

286 Time-kill synergy studies revealed RRIKA and RR, at only $\frac{1}{2} \times$ MIC enhanced the killing of
287 lysostaphin (at $\frac{1}{4} \times$ MIC) more than 1000 fold and completely eradicated MRSA and VRSA
288 isolates within 20 minutes (figure 3).

289 **Bacterial membrane disruption activity (Bacteriolysis)**

290 Since evidence of rapid cell lysis of treated cultures is indicative of membrane damaging activity
291 of antimicrobial peptides (22, 23), we assessed whether the bactericidal activity of our peptides
292 was associated with lysis of bacterial cells by measuring culture turbidity over time after
293 treatment. We monitored the turbidity of MRSA cultures treated with $4 \times$ MIC of peptides over a
294 10 hr period at OD₅₉₅ via a microplate reader. As shown in figure 4 both RRIKA and RR caused
295 a rapid decrease in OD₅₉₅ even with a high inoculum concentration of 10^8 CFU/ml. Peptide
296 RRIKA produced greater than 50% and 86% reduction in turbidity after 2.5 and 6 hr,
297 respectively. RR produced a similar percentage of turbidity as RRIKA albeit at a slower rate
298 (after 4.5 and 10 hr). Nisin at $4 \times$ MIC, a polycyclic antibacterial peptide which is known as a
299 membrane perturbing agent (24), lysed cells at a slightly higher rate than RRIKA with more than
300 70% reduction in turbidity observed after 2.5 h. In contrast, wells treated with $4 \times$ MIC of

301 vancomycin (a cell wall biosynthesis inhibitor) had no effect on culture turbidity within the same
302 time frame.

303 **Permeabilization of *S. aureus* membrane (calcein leakage assay)**

304 We monitored the effect of the peptides on MRSA membrane integrity by using the calcein
305 leakage assay as described previously (19, 25). Damage of membrane integrity is indicated by
306 calcein leakage from bacterial cells, leading to a reduction in fluorescence intensity. Figure 5
307 demonstrates that both peptides perturb the *S. aureus* cell membrane leading to leakage of
308 preloaded calcein in both a concentration- and time-dependent manner. RRIKA was faster and
309 more potent in comparison to RR in membrane perturbation. At $5 \times \text{MIC}$, RRIKA and RR
310 caused more than 60% and 30% leakage respectively within 60 minutes. When the concentration
311 of peptide was increased to $10 \times \text{MIC}$, a significant change in membrane damage was observed
312 for both peptides. There was a more than 75% and 60% reduction in fluorescence intensity
313 measured for RRIKA and RR respectively in the same time frame. Nisin at $5 \times \text{MIC}$ resulted in
314 more than 70% calcein leakage from cells, which is comparable to RRIKA at $10 \times \text{MIC}$.
315 Vancomycin, as expected, had no effect on the bacterial cell membrane integrity even at high
316 concentration ($10 \times \text{MIC}$).

317 **DNA binding properties**

318 Since there was no obvious evidence that the antibacterial effect of AMPs was restricted to
319 perforation of membranes, we explored the feasibility of other intracellular targets, such as DNA
320 by assessing the DNA-binding properties of peptides (26, 27). As shown in figure 6, both
321 peptides were able to bind the plasmid DNA and delay its electrophoretic migration into agarose
322 gel, in a dose-dependent manner. At $8 \mu\text{M}$ concentration, of RRIKA peptide, a fraction of the
323 plasmid DNA was still able to migrate into the gel as non-complexed DNA; however at $16 \mu\text{M}$,

324 the majority of DNA remained in the gel's wells. At higher concentrations, complete retardation
325 of DNA migration was exhibited, implying that the DNA was aggregated by peptide. Similar
326 patterns of migration were observed with the RR peptide except that free DNA was still seen at
327 16 μM concentration. In contrast, magainin, a cationic antimicrobial peptide of amphibian skin,
328 lacks this ability to bind DNA even at concentrations up to 80 μM .

329 **Quantification of activity against biofilms**

330 In order to evaluate the efficacy of RR and RRIKA against established biofilms, we measured
331 the biofilm mass post-treatment by crystal violet staining biofilms formed by clinical isolates of
332 *S. aureus* and *S. epidermidis*. As observed in figure 7a, both peptides significantly reduced *S.*
333 *aureus* biofilms as compared to the antibiotics of choice. RRIKA and RR, at only $2 \times \text{MIC}$,
334 dispersed more than 65% of mature biofilms ($p < 0.01$), while vancomycin and linezolid, at $16 \times$
335 MIC, was only capable of reducing 40% of biofilm mass ($p < 0.01$). There was significant
336 difference between peptide and antibiotics treated biofilms ($p < 0.01$).

337 With regards to *S. epidermidis*, both RR and RRIKA, at 32 and $64 \times \text{MIC}$, were able to reduce
338 more than 50% of biofilm mass respectively ($p < 0.01$) (figure 7b). Vancomycin and linezolid,
339 on the other hand, reduced only 9.5 and 10.7% of biofilm mass, respectively ($p < 0.01$) at
340 concentrations equivalent to $256 \times \text{MIC}$. Also we found statistical significance between peptide
341 and antibiotics treated biofilms ($p < 0.01$).

342 **Hemolysis assay and cell toxicity**

343 We assessed the release of hemoglobin from human erythrocytes exposed to different
344 concentrations of peptides. As depicted in figure 8, even at concentrations as high as 300 μM ,
345 RRIKA and RR exhibited minimal hemolysis against RBCs (maximum of 10% hemolysis

346 observed). In contrast, melittin completely lysed RBCs (100% hemolysis) at a significantly lower
347 concentration of 5 μ M.

348 The cytotoxic effect of the peptides on HeLa cells was evaluated by the MTS assay. As depicted
349 in figure 9, neither RRIKA nor RR was toxic to the mammalian cells tested up to concentrations
350 of 32 μ M and 64 μ M, respectively. These values correlate to 8 and 4 \times MIC for RRIKA and RR,
351 respectively. The half maximal effective concentration (EC_{50}) of RRIKA and RR against
352 macrophage cell lines was 64 and 128 μ M respectively. On the other hand, melittin showed high
353 toxicity even at bacteriostatic concentration with EC_{50} less than 5 μ M.

354 **DISCUSSION:**

355 In this study, we evaluated the antimicrobial activity of two short synthetic peptides, and
356 revealed their potential mechanisms of action. We observed that RRIKA and RR exhibited
357 potent antibacterial activity against all tested *S. aureus* isolates. Moreover, our peptides
358 demonstrated activity against multiple clinical isolates of MRSA, particularly MRSA USA300, a
359 community-associated strain which is responsible for outbreaks of staphylococcal skin and soft-
360 tissue infections (SSTI) in the United States (28). Similarly, the potent bactericidal activity was
361 observed in other clinical MRSA isolates (USA100, USA200 and USA500) that are resistant to
362 various antibiotic classes including macrolides, aminoglycosides, lincosamides, and
363 fluoroquinolones. Additionally, we found that these peptides retained their activity against
364 strains of *S. aureus* which have resistance to drugs of choice in treating MRSA infections,
365 namely linezolid (MRSA NRS119) and vancomycin (VRSA strains). It is worthy to note that, the
366 C terminus of the RR peptide was modified to contain three additional amino acids (isoleucine,
367 lysine and alanine) which made the RRIKA peptide more amphipathic and hydrophobic and also
368 increased its positive net charge. These modifications augmented the antibacterial activity of

369 RRIKA which exhibited MICs two to eight folds lower than those of RR. The enhanced efficacy
370 of RRIKA may reveal a functional structure activity relationship and provide a template for
371 future peptide synthesis.

372 Time-kill kinetics revealed a major advantage of the synthetic peptides over tested antibiotics.
373 While RRIKA and RR eliminated MRSA within 1-2 hr after treatment, vancomycin required 24
374 hr to reduce the bacterial CFU by 3- \log_{10} . Linezolid, on the other hand, exhibited only a
375 bacteriostatic effect.

376 Growth kinetic measurements of MRSA exposed to peptides clearly demonstrated that RRIKA
377 and RR peptides exhibited kinetic behavior similar to membrane lytic peptides such as the
378 lantibiotic nisin (24) and amphibian magainin (23) but different from non-lytic peptides such as
379 fungal plectasin (29) and amphibian buforin II (23). These data suggest that our peptides act by
380 disrupting the integrity of the bacterial membrane. To validate this hypothesis, we studied the
381 effect of peptides on MRSA membranes using a calcein leakage assay. Similar to the well-
382 known membrane damaging peptide, nisin, both RRIKA and RR permeabilized *S. aureus* cells in
383 a concentration and time-dependent manner. These observations clearly validate that the
384 mechanism of bacterial killing by our peptides is mediated by pore formation and disruption of
385 the bacterial cell membrane, leading to leakage of cytoplasmic contents and ultimately cell lysis.
386 However, recent evidence suggests that lysis of bacterial cells by pore formation is not the only
387 mechanism of microbial killing of some AMPs and there are other possible intracellular targets,
388 such as aggregation of DNA (26). To further explore this possibility, we examined the bacterial
389 DNA-binding ability of the peptides by electrophoretic gel retardation. Our data indicated that
390 both RRIKA and RR bind with bacterial DNA and alter its electrophoretic mobility in agarose
391 gels. Although a detailed study on the interaction between peptides and nucleic acids *in vivo* is

392 needed, the current results lead us to suggest that RRIKA and RR might inhibit the bacterial
393 functions by binding to bacterial DNA after perturbing the cell membranes, resulting in an
394 augmented effect by two different mechanisms. While RRIKA and RR are not toxic to
395 mammalian cells at the levels used here, possible crossing of the peptides to the nuclear
396 membrane and binding to eukaryotic DNA must be examined further.

397 Major limitations with the use of AMPs for systemic applications are their possible inactivation
398 by serum or physiological concentrations of salts. Serum inactivate peptides either through
399 cleavage by proteases or through binding with proteins or lipids (30) We observed that RRIKA
400 but not RR retained its bactericidal activity, although at a slower rate, in the presence of 10%
401 FBS. Furthermore, both RRIKA and RR retain their antibacterial activity when tested in
402 increasing concentration of NaCl and MgCl₂; in contrast to many well studied AMPs (such as
403 LL-37, human β -defensin-1, gramicidins, bactenecins, and magainins) which showed
404 substantially reduced antibacterial activities under the same conditions (11). Previously, Turner
405 *et al.* reported that LL-37 and HNP-1 demonstrated a 12-fold and 100-fold increase in the MIC
406 of MRSA, respectively when 100 mM NaCl was added to the test medium (10). The ability to
407 resist the effects of salt and serum provide a selective advantage for our peptides for potential
408 therapeutics in physiological solutions.

409 After identifying that our peptides were capable of inhibiting bacterial growth alone, we wanted
410 to explore their ability to be used in combination with other antimicrobials such as lysostaphin.
411 Lysostaphin is a zinc metalloenzyme that specifically cleaves the abundant pentaglycine cross-
412 bridges of the staphylococcal cell wall. Several studies have reported the potential applications
413 of lysostaphin in the treatment of staphylococcal infections, however, *S. aureus* has developed
414 resistance against lysostaphin via different mechanisms (31). When we tested both RR and

415 RRIKA in combination with lysostaphin, we observed a synergistic relationship demonstrated by
416 complete eradication of MRSA and VRSA within 20 minutes (at very low concentrations). The
417 observed synergistic effect may be due the cleaving of the cell wall peptidoglycan by
418 lysostaphin, which allows more access of peptides to the bacterial membrane. The synergistic
419 relationship observed between our peptides and lysostaphin is important as it has the advantage
420 of reducing the emergence of bacterial resistance to both agents while also minimizing drug-
421 associated toxicity (by lowering the therapeutic dose needed for each antimicrobial agent to
422 effectively treat MRSA infection).

423 One of the difficult challenges facing current antibiotics is bacterial biofilms, which are vital in
424 the pathogenesis of staphylococcal infections. Biofilms hinder the penetration of antimicrobials
425 to access bacteria, leading to failure of treatments (32). Our data showed that both RR and
426 RRIKA are capable of disrupting *S. aureus* and *S. epidermidis* biofilms more efficiently than
427 drugs of choice (vancomycin and linezolid). Most bacteria living in biofilms are either slow-
428 growing or non-growing dormant cells that are difficult to treat with antibiotics that normally
429 inhibit macromolecular synthesis in growing cells (33). However our peptides target mainly the
430 microbial cell membrane, a characteristic that is not only in dividing organisms, but also in
431 quiescent cells or stationary phase bacteria (34).

432 Cell toxicity is one of the major limitations in the development of antibacterials, particularly if
433 the target of action is the cell membrane (7). We observed minimal hemolysis (less than 10%)
434 with our peptides even with high concentrations (300 μ M). On the other hand, 5 μ M of melittin
435 completely lysed human RBCs (100% hemolysis). Furthermore, RRIKA and RR were not toxic
436 to HeLa cells at 8 and 4 \times MIC, respectively.

437 In conclusion, our findings reveal the potent bactericidal action of peptides RRIKA and RR
438 against MRSA. The mechanism of action of the peptides is mainly due to bacterial lysis as a
439 consequence of bacterial membrane disruption and possibly by binding of the peptides to
440 bacterial DNA, which interferes with necessary cellular functions vital for microbial survival.
441 Such effects are extremely challenging for pathogens to overcome by developing resistance;
442 unlike current antibiotics, which usually inhibit metabolic pathways that can lead to bacterial
443 resistance (8). To date, issues such as poor pharmacokinetics have limited the potential systemic
444 applications of therapeutic AMPs (7, 8, 30). Therefore, AMPs which have advanced into
445 preclinical or clinical trials, are indicated for topical treatment of bacterial skin infections (7, 8,
446 30) such as Pexiganan (Access Pharmaceuticals, Inc.) for curing diabetic foot ulcers (35). *S.*
447 *aureus* is the most frequently isolated microorganism in diabetic foot infections, and those
448 caused by MRSA are associated with the worse outcomes and more frequent amputations (36-
449 41). Additionally, bacterial biofilms appear to play an important role in increasing the difficulty
450 of treating these ulcers (40). Furthermore, it has been proven recently that the clinical severity of
451 *S. aureus* skin infection is driven by the inflammatory response to the bacteria, rather than
452 bacterial burden (42-44). Taken together, the characteristics of the presented peptides with
453 combined bactericidal, anti-biofilm and anti-inflammatory effect may offer an effective way for
454 treating Staphylococcal skin infection. Therefore, these results support the potential for further
455 study and development of RRIKA and RR as topical therapeutics particularly in an era of
456 emerging drug resistance.

457 **ACKNOWLEDGEMENT**

458 Mohamed F. Mohamed and Maha I. Hamed are supported by a scholarship from the Egyptian
459 Cultural and Educational Bureau (ECEB) in Washington DC, USA. We would like to thank the

460 Network of Antimicrobial Resistance in *Staphylococcus aureus* (NARSA) program supported
461 under NIAID/NIH Contract # HHSN272200700055C for providing *Staphylococcus aureus*
462 strains used in this study. The authors wish to acknowledge Mr Haroon Mohammed for editing
463 the manuscript.

464

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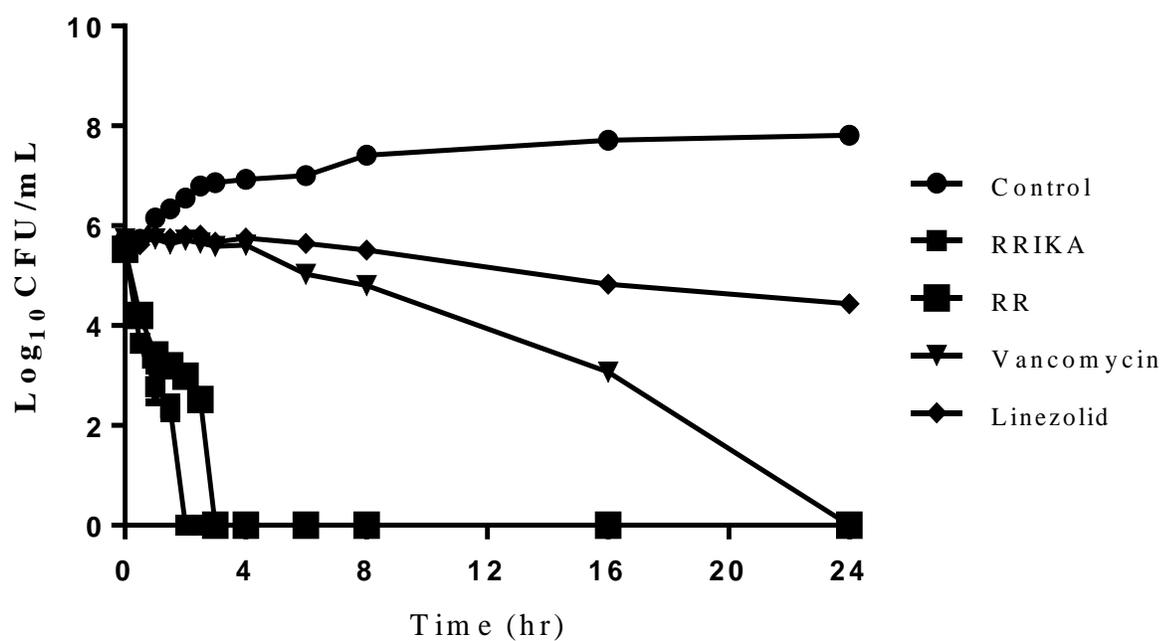


Figure 1: Bacterial killing kinetics of RRIKA, RR, vancomycin and linezolid at 5X MIC against MRSA USA300 in MHB (Mueller Hinton broth). Samples treated with peptide diluent (sterile water) were used as a control. The results are given as means \pm SD (n = 3; data without error bars indicate that the SD is too small to be seen)

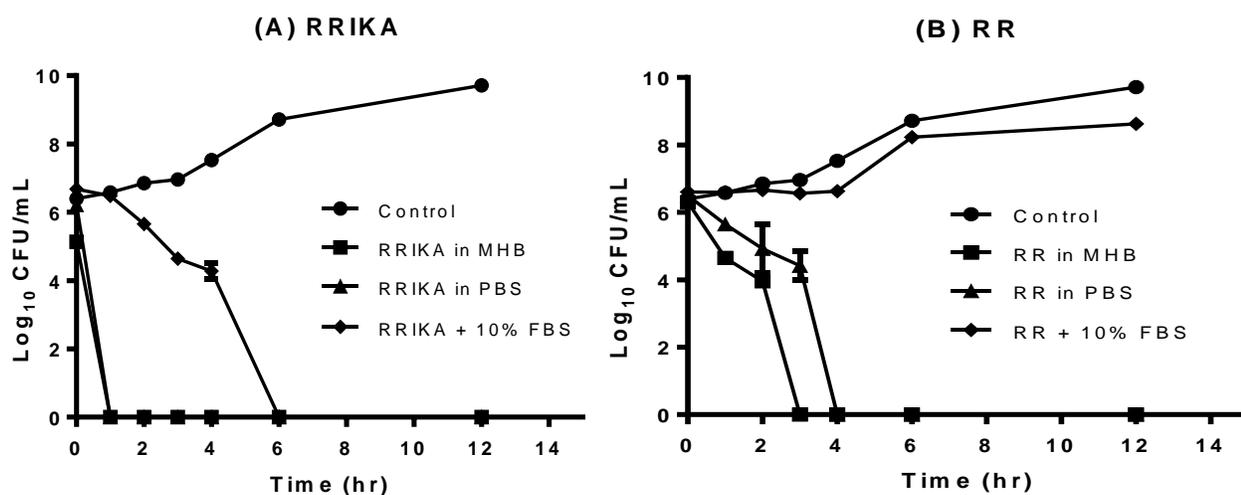


Figure 2: Effect of culture conditions on killing kinetics of RRIKA (A), RR (B) at 5X MIC against MRSA USA300. Abbreviations, MHB “Mueller Hinton broth”, PBS “phosphate buffered saline” or MHB+10% FBS “fetal bovine serum”. Untreated samples were used as a control was. The killing curves were identical (overlapping in the figure) for RRIKA in MHB and in PBS. Each experiment was done in triplicate, and the values represent means \pm SD (data without error bars indicate that the SD is too small to be seen).

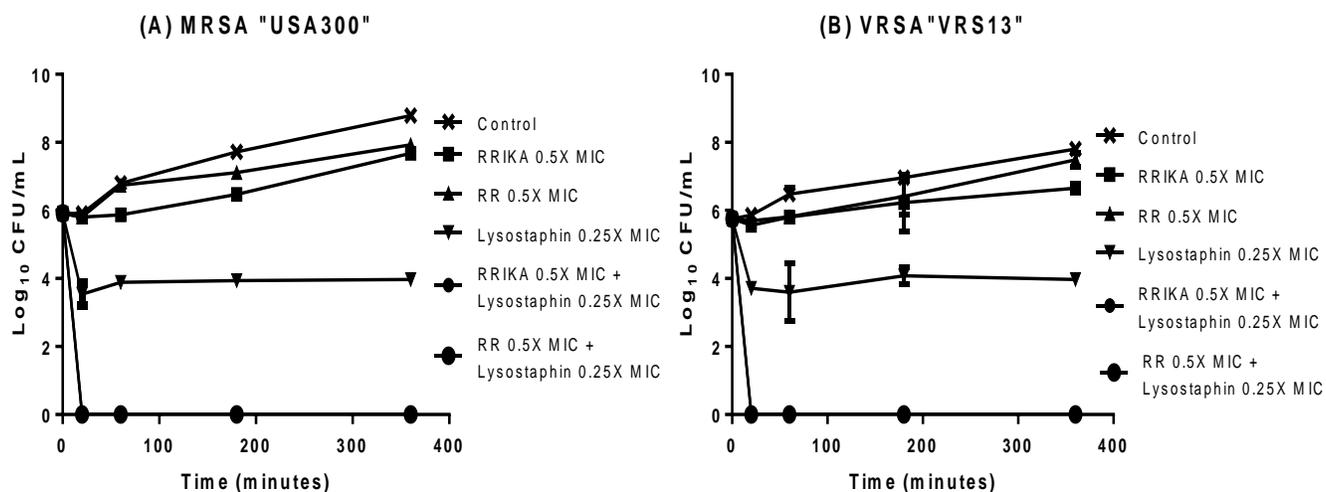


Figure 3: Synergistic killing of RRIKA and RR with lysostaphin by time kill assay. MRSA USA300 (A) and VRSA VRS13 (B) were incubated with 0.5X MIC of RRIKA or RR alone or in combination with 0.25X MIC of lysostaphin. Samples were obtained at different time points, then plated and counted. Control was untreated samples. The killing curves were identical (overlapping in the figure) for RRIKA and RR in combination with lysostaphin. The results are given as means \pm SD (n = 3; data without error bars indicate that the SD is too small to be seen)

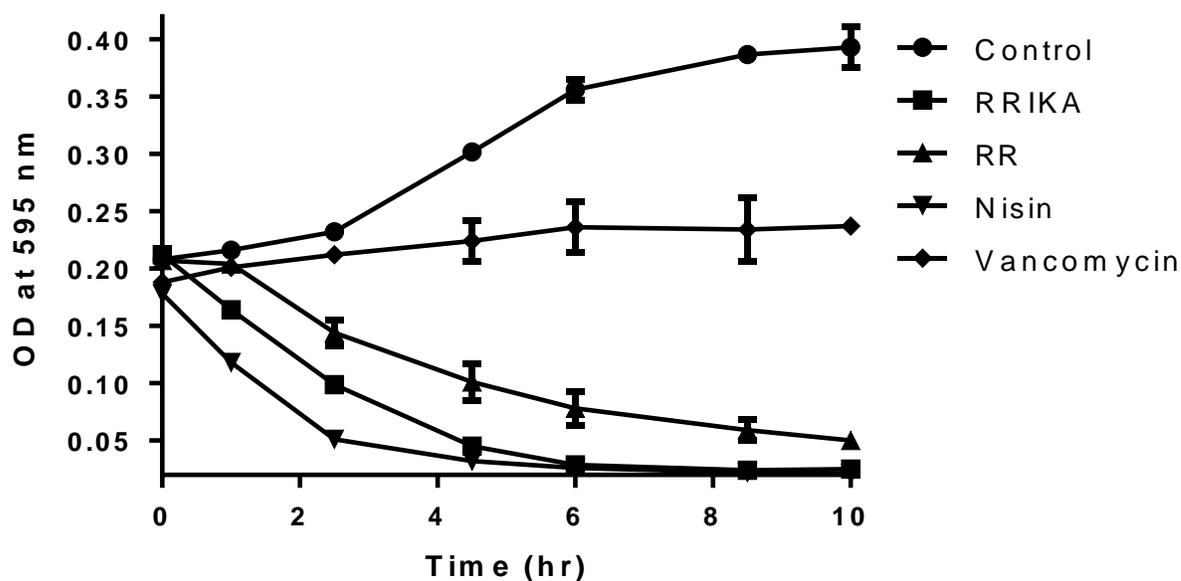


Figure 4: Killing kinetics of MRSA USA300 exposed to 4X MIC of RRIKA, RR, nisin, vancomycin and sterile water (control) as observed by measuring OD595 by microplate reader over time. Results are representative of two separate experiments, each was done in triplicate. Error bars represent standard deviation values (data without error bars indicate that the SD is too small to be seen)

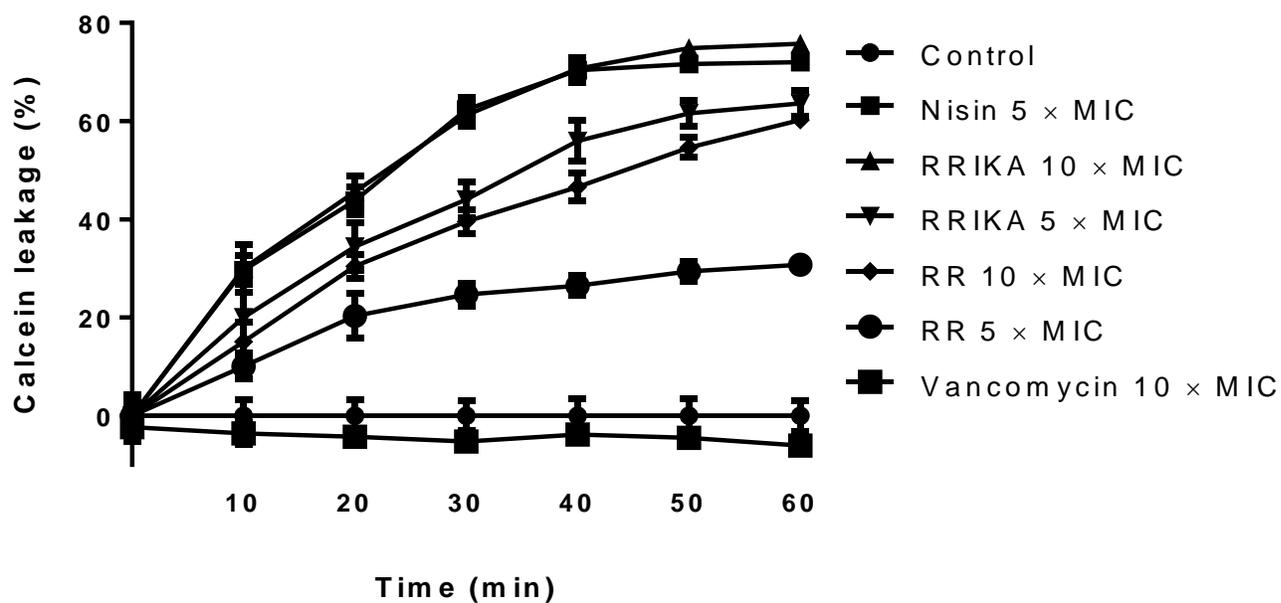


Figure 5: Permeabilization of the cytoplasmic membrane of MRSA USA300 as a function of peptide concentration, indicated by percent of calcein leakage for 60 min exposure. The results are given as means \pm SD ($n = 3$; data without error bars indicate that the SD is too small to be seen).

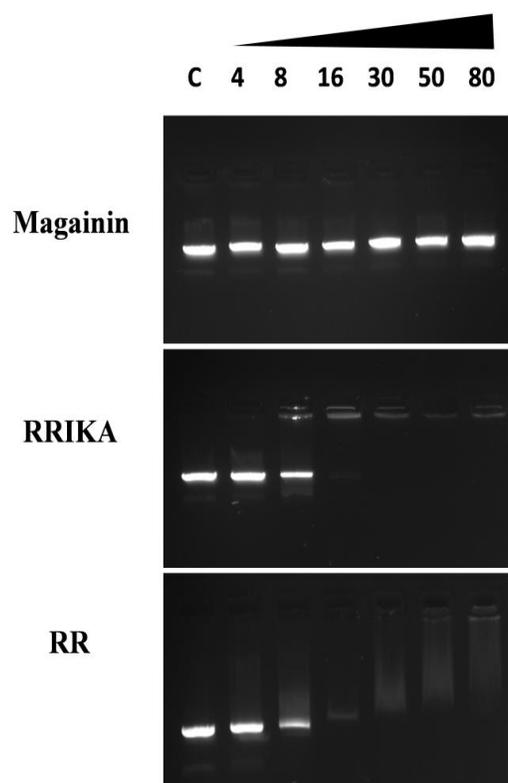


Figure 6: Interaction of peptides with plasmid DNA. Binding was assayed by measuring inhibition of migration by plasmid DNA. Different concentrations of peptides were incubated with 250 ng for 1 hr at room temperature prior to electrophoresis on a 1.0% agarose gel. The numbers above the lanes represent the concentration (in μM) of magainin, RRIKA or RR. Lane C, control consisting of plasmid DNA only.

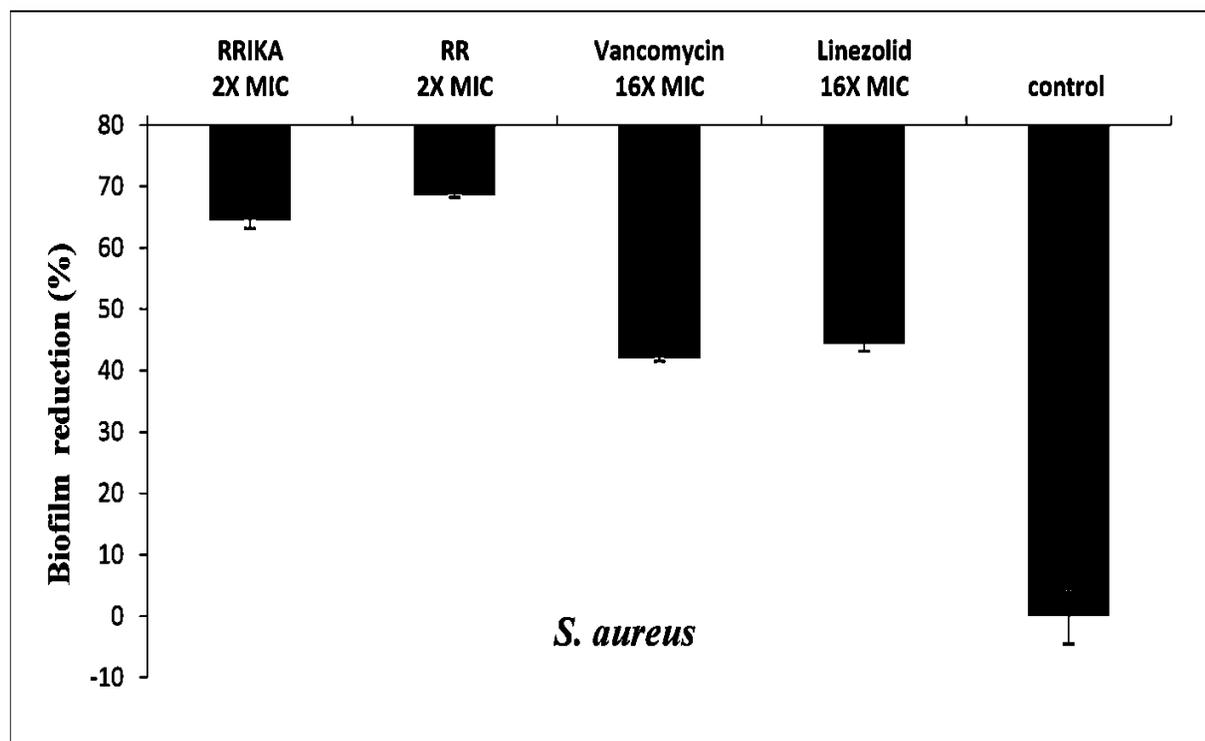


Fig. 7a

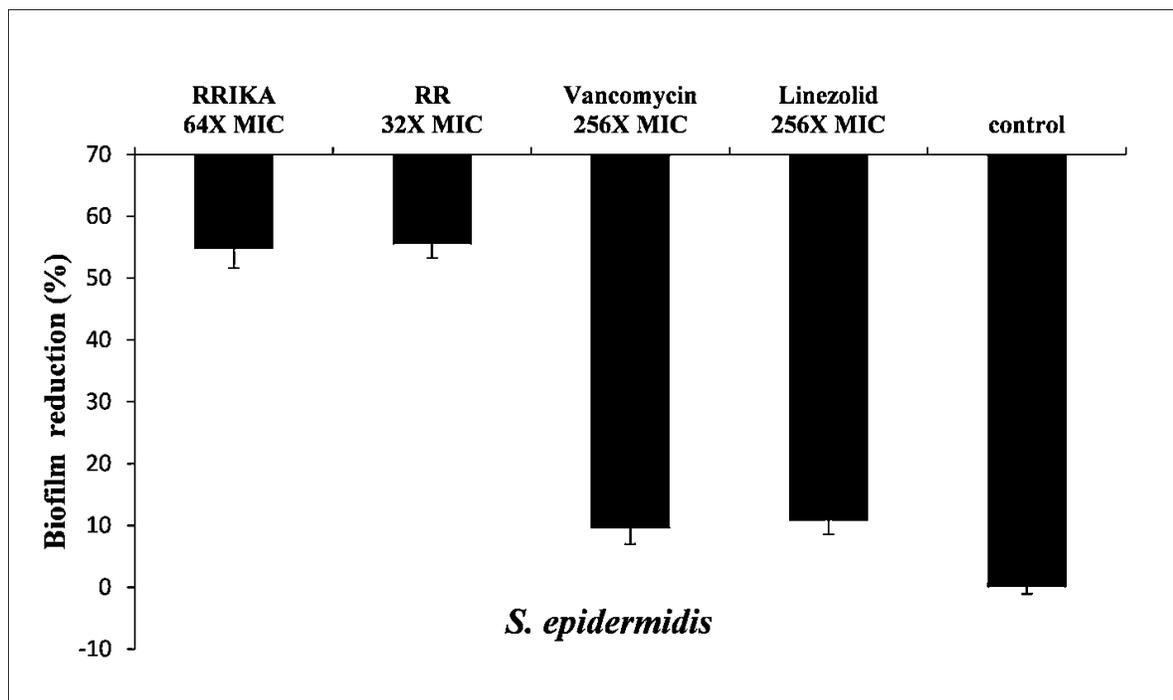


Fig. 7b

Figure 7: The effect of peptides (RRIKA & RR) and antibiotics (vancomycin & linezolid) on established biofilms of *S. aureus* (a) and *S. epidermidis* (b). The adherent biofilm stained by crystal violet, then the dye was extracted with ethanol, measured at 595 nm absorbance and presented as percentage of biofilm reduction compared to untreated wells “control”. All experiments were done in triplicate for statistical significance. One asterisk (*) indicates statistically different than the positive control ($p < 0.01$). Two asterisks (**) indicates statistically different than the antibiotic treated wells ($p < 0.01$).

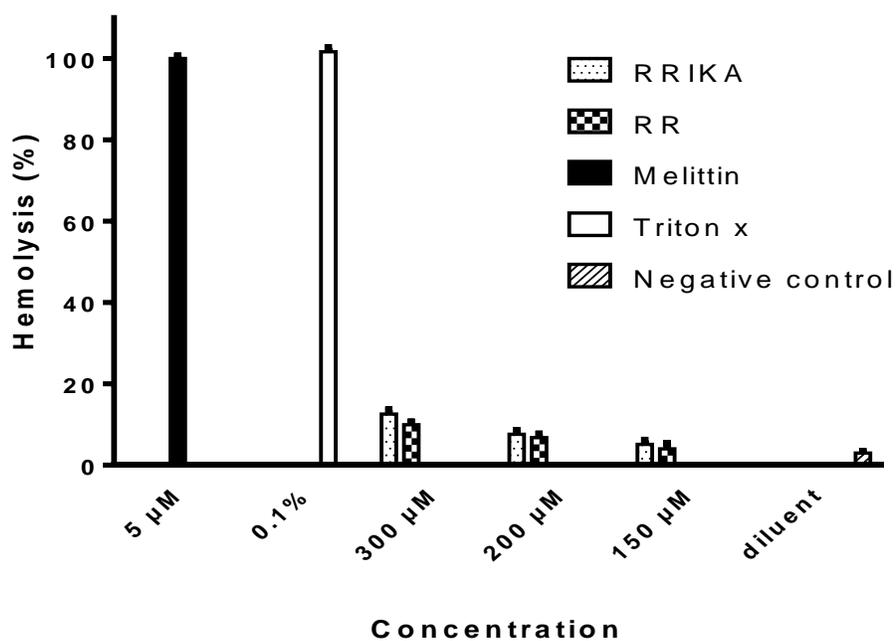


Figure 8: The release of hemoglobin in the supernatant of human erythrocytes after treatment with increasing amounts of RRIKA and RR was measured at 415 nm. Data collected after 1 hr of incubation are presented. Melittin (5 μ M) and 0.1 % of triton X-100 served as positive controls. Phosphate buffered saline (PBS) served as a negative control.

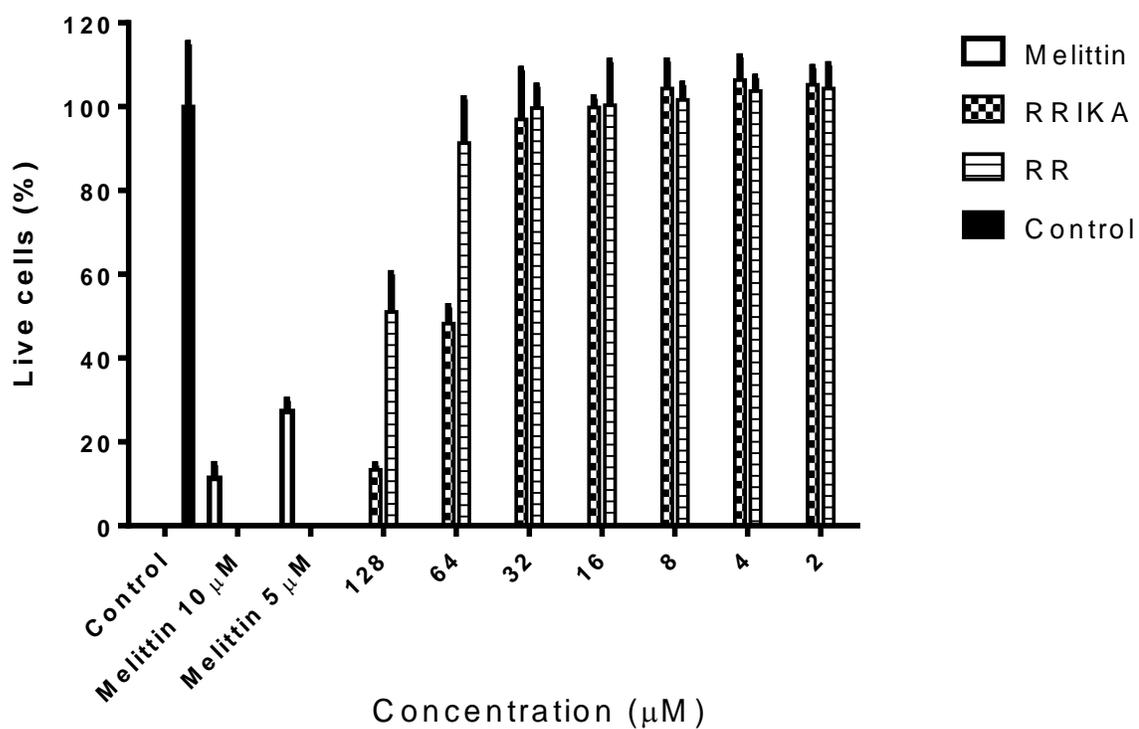


Figure 9: Cytotoxicity assay showing the percent mean absorbance at 490 nm after incubating HeLa cells with RRIKA and RR and melittin at different concentrations. Diluent was used as a control. Cell viability was measured by MTS assay. Results are expressed as means from three measurements \pm standard deviation.

Table 1: Minimum inhibitory concentration (MIC) of peptides and antibiotics against clinical and drug-resistant *Staphylococci* strains

Strain Type	Strain ID	Origin	Phenotypic Properties	MIC(μ M)					
				Peptide RRIKA	Peptide RR	Lin	Van	Nisin	Amp
Methicillin sensitive <i>Staphylococcus aureus</i> (MSSA)	ATCC 6538	-	Quality control and biofilm forming strain	2	16	2	0.5	1	≤ 0.25
	RN4220	United States	Resistant to mupirocin	4	8	1	0.5	nd	≤ 0.25
	NRS72	United Kingdom	Resistant to penicillin	4	32	4	0.5	2	>32
	NRS77	United Kingdom	-	4	16	2	0.5	2	≤ 0.25
	NRS846	-	-	4	16	4	0.5	2	≤ 0.25
NRS860	-	-	-	4	16	1	0.5	2	≤ 0.25
Methicillin resistant <i>Staphylococcus aureus</i> (MRSA)	USA100	United States (Ohio)	Resistant to ciprofloxacin, clindamycin, erythromycin	2	16	4	0.5	nd	>32
	USA200	United States (North Carolina)	Resistant to ciprofloxacin, clindamycin, erythromycin, gentamicin, and methicillin	4	8	4	0.5	nd	>32
	USA300	United States (Mississippi)	Resistant to erythromycin, methicillin, and tetracycline	4	16	4	0.5	1	>32
	USA400	United States (North Dakota)	Resistant to methicillin and tetracycline	2	8	4	0.5	nd	>32
	USA500	United States	Resistant to ciprofloxacin,	4	8	4	1	nd	>32

		(Connecticut)	clindamycin, erythromycin, gentamicin, methicillin, tetracycline, and trimethoprim						
	USA700	United States (Louisiana)	Resistant to erythromycin and methicillin	2	8	4	1	nd	>32
	USA800	United States (Washington)	Resistant to methicillin	2	8	4	0.5	nd	>32
	USA1000	United States (Vermont)	Resistant to erythromycin and methicillin	4	8	4	0.5	nd	>32
	USA1100	United States (Alaska)	Resistant to methicillin	2	8	4	1	nd	>32
	NRS194	United States (North Dakota)	Resistant to methicillin	2	8	4	1	nd	>32
	NRS108	France	Resistant to gentamicin	4	8	4	0.5	nd	>32
	NRS119 (Lin ^R)	United States (Massachusetts)	Resistant to linezolid	4	16	128	0.5	nd	>32
	ATCC 43300	United States (Kansas)	Resistant to methicillin	4	16	4	0.5	nd	>32
	ATCC BAA- 44	Lisbon, Portugal	Multidrug-resistant strain	4	16	4	0.5	nd	>32
	NRS70	Japan	Resistant to erythromycin and spectinomycin	4	16	2	0.5	2	>32
	NRS71	United Kingdom	Resistant to tetracycline and methicillin	4	16	2	0.5	2	>32
	NRS100	United States	Resistant to tetracycline and methicillin	4	16	2	0.5	2	>32
	NRS123	United States (North Dakota)	Resistant to tetracycline and methicillin	4	16	2	0.5	1	>32
Vancomycin intermediate <i>Staphylococcus aureus</i>	NRS1	Japan	Resistant to aminoglycosides and tetracycline Glycopeptide-	2	16	2	8	nd	>32

(VISA)			intermediate <i>Staphylococcus aureus</i>						
	NRS19	United States (Illinois)	Glycopeptide- intermediate <i>Staphylococcus aureus</i>	4	8	2	4	nd	>32
	NRS37	France	Glycopeptide- intermediate <i>Staphylococcus aureus</i>	2	8	2	8	nd	>32
Vancomycin Resistant <i>Staphylococcus aureus</i> (VRSa)	VRS1	United States	Resistant to vancomycin	4	32	1	>128	1	>32
	VRS2	United States	Resistant to vancomycin	4	16	1	16	1	>32
	VRS3a	United States	Resistant to vancomycin	4	16	1	128	2	>32
	VRS4	United States	Resistant to vancomycin	4	32	2	128	nd	>32
	VRS5	United States	Resistant to vancomycin	4	32	2	128	nd	>32
	VRS6	United States	Resistant to vancomycin	4	32	1	>128	nd	2
	VRS7	United States	Resistant to vancomycin	4	16	2	128	nd	>32
	VRS8	United States	Resistant to vancomycin	4	16	1	>128	nd	32
	VRS9	United States	Resistant to vancomycin	4	16	1	>128	nd	>32
	VRS10	United States	Resistant to vancomycin	4	16	4	>128	nd	>32
	VRS11b	United States	Resistant to vancomycin	4	32	1	>128	nd	>32
	VRS12	United States	Resistant to vancomycin	4	32	2	>128	nd	>32
VRS13	United States	Resistant to vancomycin	4	16	1	>128	1	16	
<i>S. epidermidis</i>	NRS101	United States	Prototype biofilm producer, Resistant to methicillin and gentamicin	2	8	2	0.5	nd	>32

Lin: linezolid, Van: vancomycin, Amp: ampicillin, nd: not determined

Table 2: Amino acid sequence and physicochemical properties of peptides used in this study

Peptide	Amino acid sequence	Length	Molecular weight	Charge	% of amino acids that are hydrophobic
RR	WLRRIKAWLRR	10	1553.9	+ 5	54 %
RRIKA	WLRRIKAWLRRIKA	13	1866.3	+ 6	57 %
KAF	KAFAKLAARKA	11	1174.4	+ 4	63 %
FAK	FAKLAARLYRKA	12	1407.7	+ 4	58 %

Table 3: Fractional inhibitory concentration (FIC) index for the combination of peptides together or with lysostaphin

Compound	FIC index ^a					
	MSSA (RN4220)		MRSA (USA300)		VRSA (VRS13)	
	RRIKA	RR	RRIKA	RR	RRIKA	RR
RRIKA	-	0.75	-	0.75	-	0.75
RR	0.75	-	0.75	-	0.75	-
Lysostaphin	0.26	0.26	0.26	0.26	0.26	0.26

^a The FIC index was determined in the presence of a fixed concentration of peptide, equivalent to $\frac{1}{4} \times \text{MIC}$.