

An antimicrobial peptide as a potential therapy for bacterial pneumonia that alleviates antimicrobial resistance

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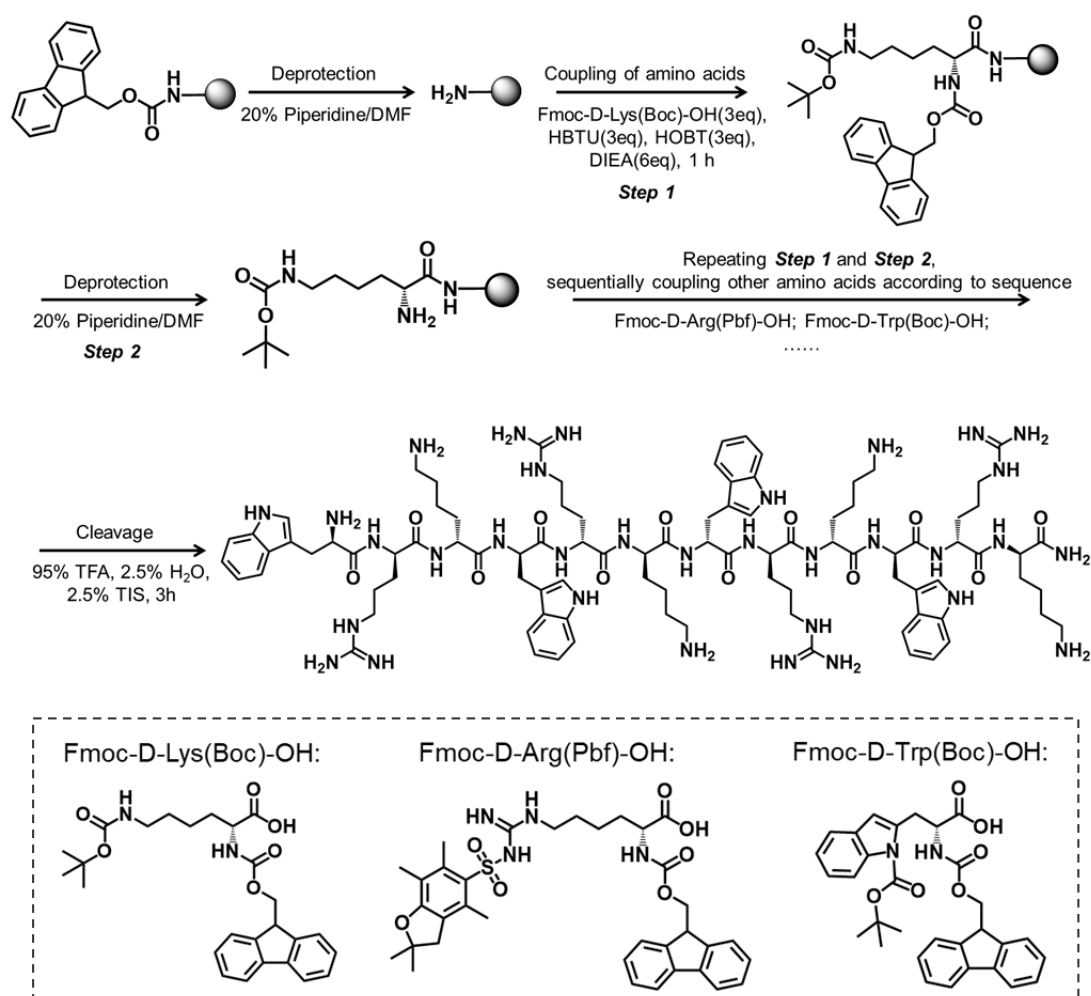
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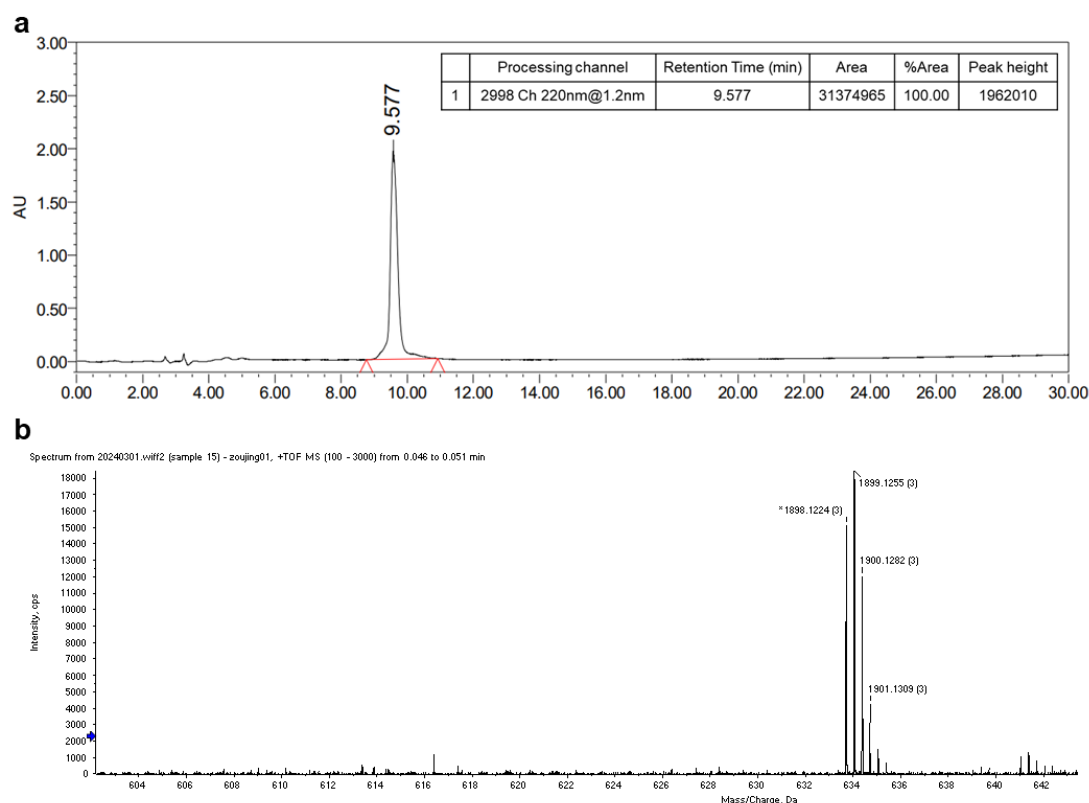
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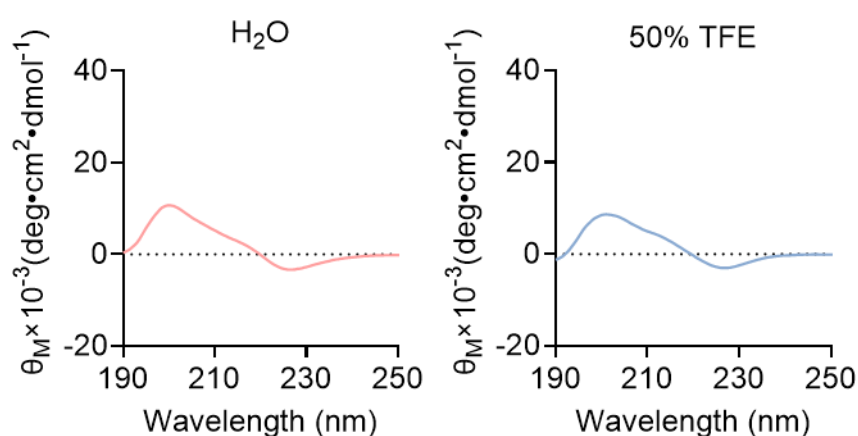
Supplementary Figures



Supplementary Fig. 1 The synthetic scheme of AMP 1003 and the amino acid structure used in the synthesis process. All amino acids are in the D-configuration unless otherwise specified.



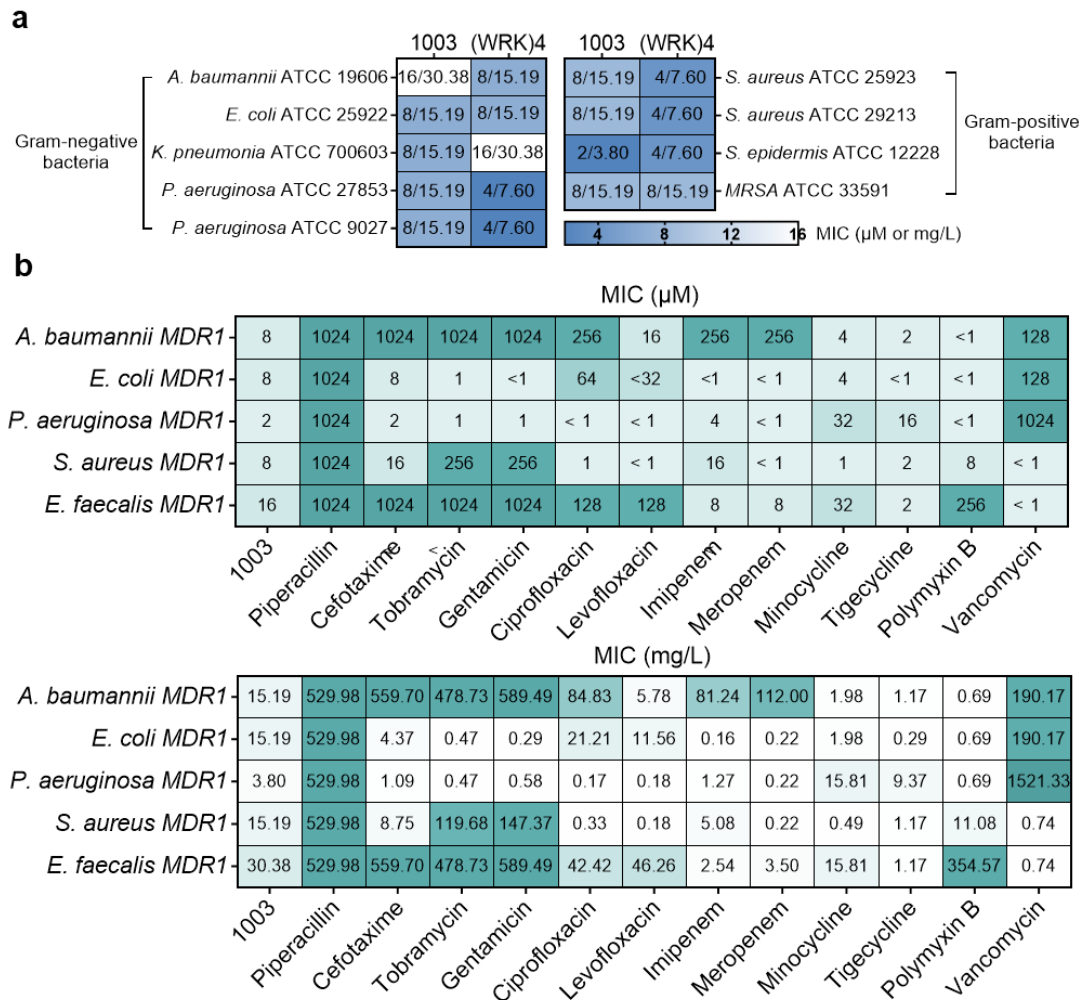
Supplementary Fig. 2 a Purity of AMP 1003 analyzed by RP-HPLC on a C18 column using a gradient of buffer B (0.1% TFA in CH₃CN/H₂O 95:5 v/v) in buffer A (0.1 %TFA in CH₃CN/H₂O 5:95 v/v) for 30 min with a flow rate of 1.0 mL/min. **b** Mass spectrum of AMP 1003 determined by EI-MS.



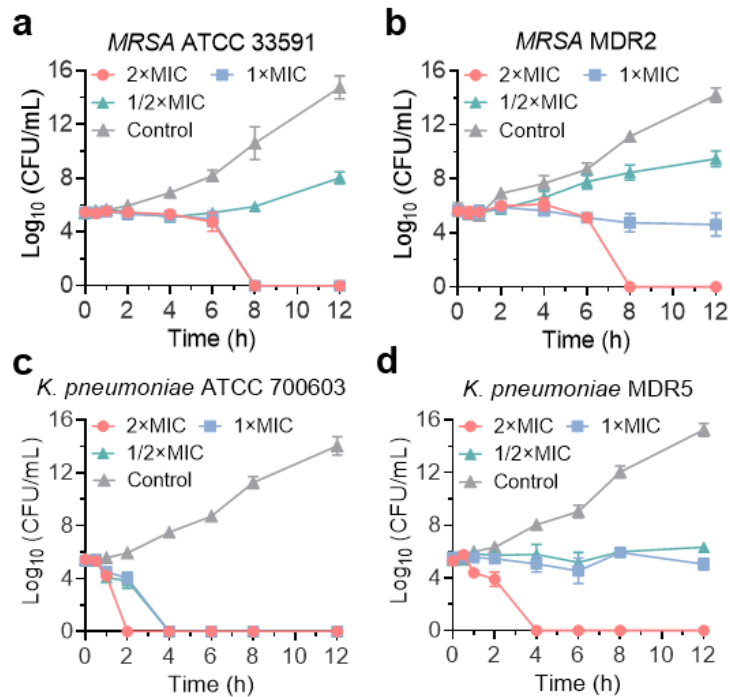
Supplementary Fig. 3 Circular dichroism (CD) spectrum of AMP 1003 in H₂O or 50% TFE. The mean residual ellipticity was employing by the formula: $\theta_M = (\theta_{\text{obs}} \times 1000) / (c \times l \times n)$. Where θ_M is the residue ellipticity (deg·cm²·dmol⁻¹), θ_{obs} is the observed ellipticity corrected for the buffer at a given wavelength (mdeg), c is the peptide concentration (mM), l is the path length (mm), and n is the number of amino acids.

a													
<i>MRSA</i> ATCC 33591	15.19	529.98	559.70	29.92	147.37	42.42	46.26	20.31	14.00	3.95	0.29	44.32	0.74
<i>MRSA</i> MDR1	15.19	33.12	8.75	0.23	0.29	0.08	0.18	0.16	0.22	0.25	0.29	2.77	0.74
<i>MRSA</i> MDR2	15.19	1059.96	559.70	59.84	147.37	678.64	46.26	81.24	3.50	3.95	1.17	22.16	0.74
<i>MRSA</i> MDR3	15.19	66.25	17.49	0.23	0.29	0.17	0.18	2.54	0.22	0.49	0.29	44.32	0.74
<i>MRSA</i> MDR4	15.19	1059.96	559.70	59.84	147.37	678.64	46.26	81.24	7.00	7.90	0.59	22.16	0.74
<i>MRSA</i> MDR5	15.19	33.12	139.92	29.92	73.69	0.33	0.18	2.54	0.22	0.25	0.29	44.32	0.74
<i>MRSA</i> MDR6	7.60	132.50	34.98	0.23	0.29	1.33	0.18	10.16	0.44	0.49	0.29	11.08	0.74
	1003	Piperacillin	Cefotaxime	Tobramycin	Gentamicin	Ciprofloxacin	Levofloxacin	Imipenem	Meropenem	Minocycline	Tigecycline	Polymyxin B	Vancomycin
b													
<i>K. pneumoniae</i> ATCC 700603	15.19	66.25	139.92	3.74	9.21	0.33	1.45	1.27	0.22	0.49	1.17	1.39	380.33
<i>K. pneumoniae</i> MDR1	121.54	1059.96	34.98	7.48	36.84	84.83	5.78	10.16	0.22	3.95	1.17	5.54	380.33
<i>K. pneumoniae</i> MDR2	243.07	1059.96	1119.40	29.92	73.69	169.66	23.13	5.08	14.00	63.22	1.17	1.39	1521.33
<i>K. pneumoniae</i> MDR3	243.07	1059.96	1119.40	29.92	73.69	84.83	46.26	20.31	28.00	126.45	1.17	0.69	760.66
<i>K. pneumoniae</i> MDR4	243.07	1059.96	1119.40	957.46	1178.97	42.42	23.13	1.27	7.00	31.61	0.59	1.39	760.66
<i>K. pneumoniae</i> MDR5	30.38	1059.96	559.70	0.47	0.58	339.32	185.02	81.24	224.01	126.45	18.74	22.16	760.66
	1003	Piperacillin	Cefotaxime	Tobramycin	Gentamicin	Ciprofloxacin	Levofloxacin	Imipenem	Meropenem	Minocycline	Tigecycline	Polymyxin B	Vancomycin

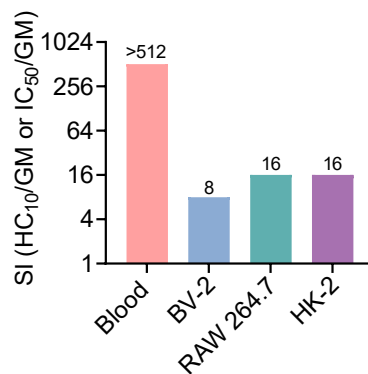
Supplementary Fig. 4 a-b Antimicrobial activity of AMP 1003 against standard and clinical isolates of multidrug-resistant *MRSA* (a) or *K. pneumoniae* (b), represented by minimum inhibitory concentration (MIC (mg/L)).



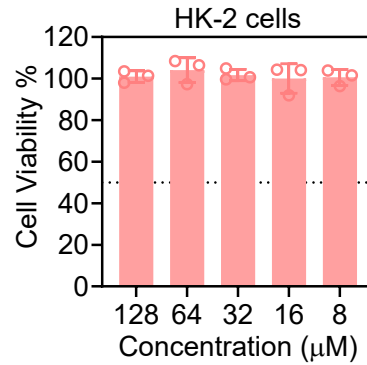
Supplementary Fig. 5 a MICs (μM or mg/L) of AMP 1003 and (WRK)₄ against standard bacteria. **b** MICs (μM or mg/L) of AMP 1003 and antibiotics against clinically isolated multidrug-resistant bacteria.



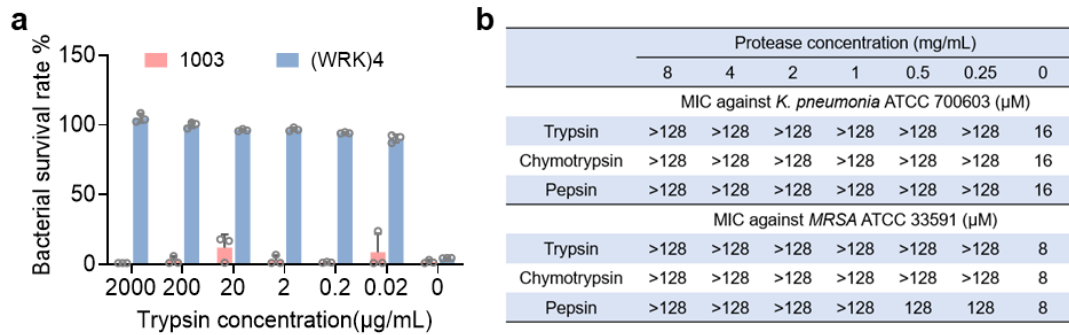
Supplementary Fig. 6 **a** and **b** Killing kinetics of vancomycin against *MRSA* ATCC 33591 (**a**) and *MRSA* MDR2 (**b**). **c** and **d** Killing kinetics of polymyxin B against *K. pneumoniae* ATCC 700603 (**c**) and *K. pneumoniae* MDR5 (**d**), $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the median, and the error bars indicate the range.



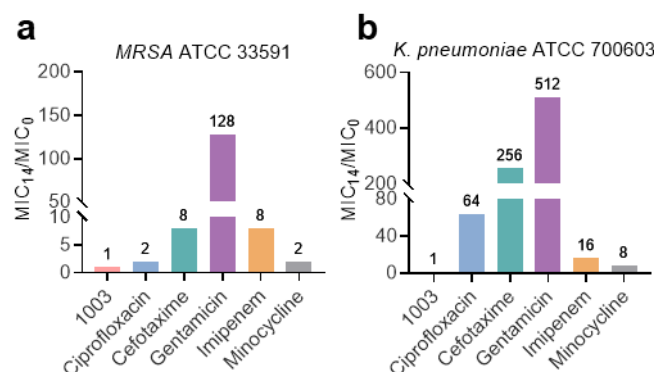
Supplementary Fig. 7 Selectivity index (SI) of AMP 1003 towards blood cells, BV-2 cells, RAW 264.7 cells and HK-2 cells.



Supplementary Fig. 8 Cytotoxicity of polymyxin B for HK-2 cells. $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.

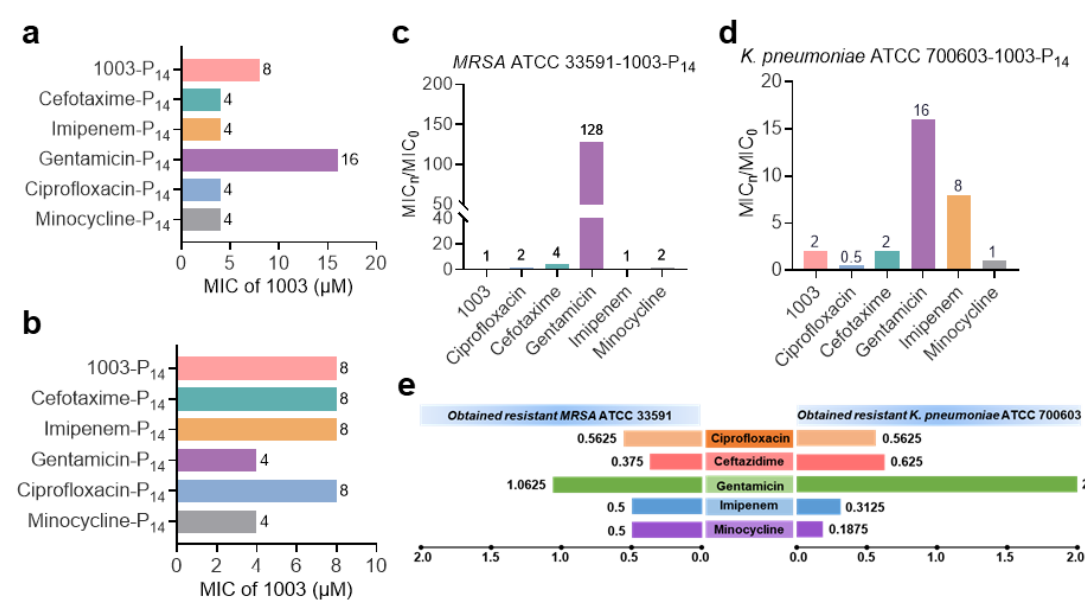


Supplementary Fig. 9 a The bacterial survival rate of AMP 1003 and (WRK)4 against *MRSA* ATCC 33591 is detected after incubation with different concentrations of trypsin for 6 h, $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. **b** MICs of (WRK)4 against *K. pneumoniae* ATCC 700603 and *MRSA* ATCC 33591 after incubation with different concentrations of trypsin, chymotrypsin or pepsin for 6 h.

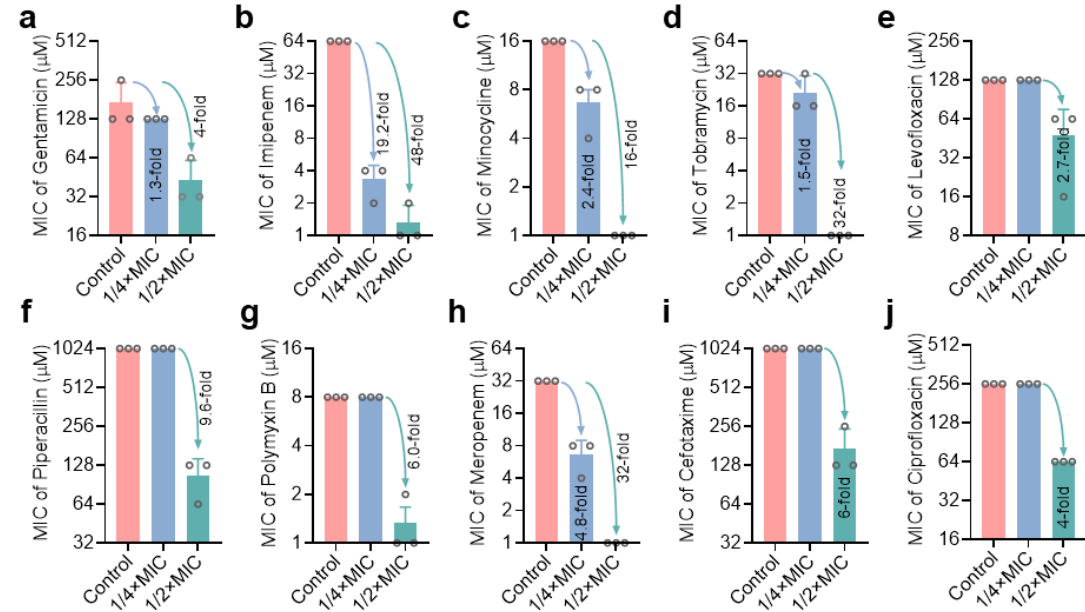


Supplementary Fig. 10 a The changes in MIC values of AMP 1003 and antibiotics against *MRSA* ATCC 33591 after treating with 1003 or antibiotics continuously at $1/2 \times \text{MIC}$ for 14 passages compared to that for 0 passage. **b** The changes in MIC values of AMP 1003 and

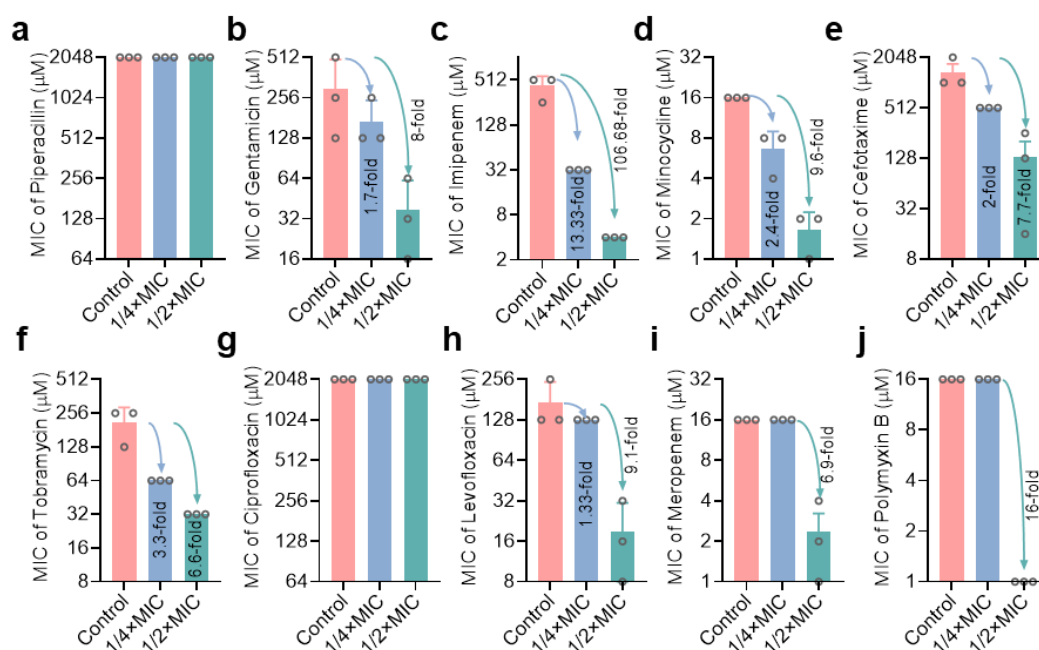
antibiotics against *K. pneumoniae* ATCC 700603 after treating with 1003 or antibiotics continuously at $1/2 \times \text{MIC}$ for 14 passages compared to that for 0 passage.



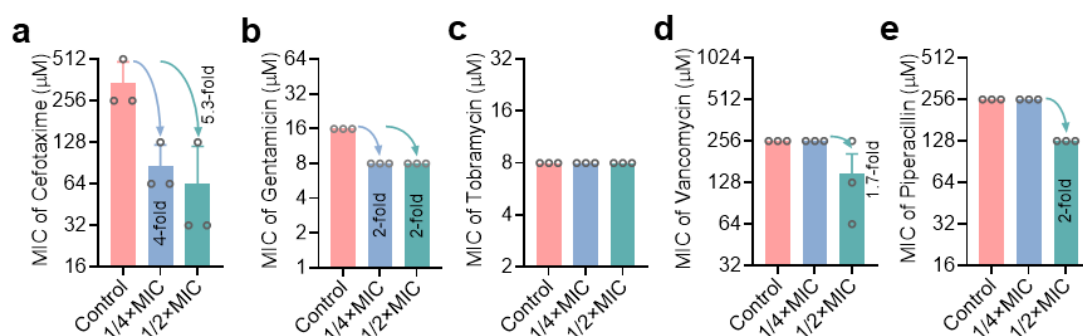
Supplementary Fig. 11 **a** and **b** The MIC values of AMP 1003 against the final-generation *MRSA* ATCC 33591 (**a**) and *K. pneumoniae* ATCC 700603 (**b**) after treating with 1003 or antibiotics continuously at $1/2 \times \text{MIC}$ for 14 passages. **c** and **d** The changes in MIC values of AMP 1003 and antibiotics against the final-generation *MRSA* ATCC 33591 (**c**) and *K. pneumoniae* ATCC 700603 (**d**) after treating with 1003 continuously at $1/2 \times \text{MIC}$ for 14 passages compared to that for 0 passage. **e** The FICIs of AMP 1003 in combination with antibiotics against the obtained resistant *MRSA* ATCC 33591 and *K. pneumoniae* ATCC 700603.



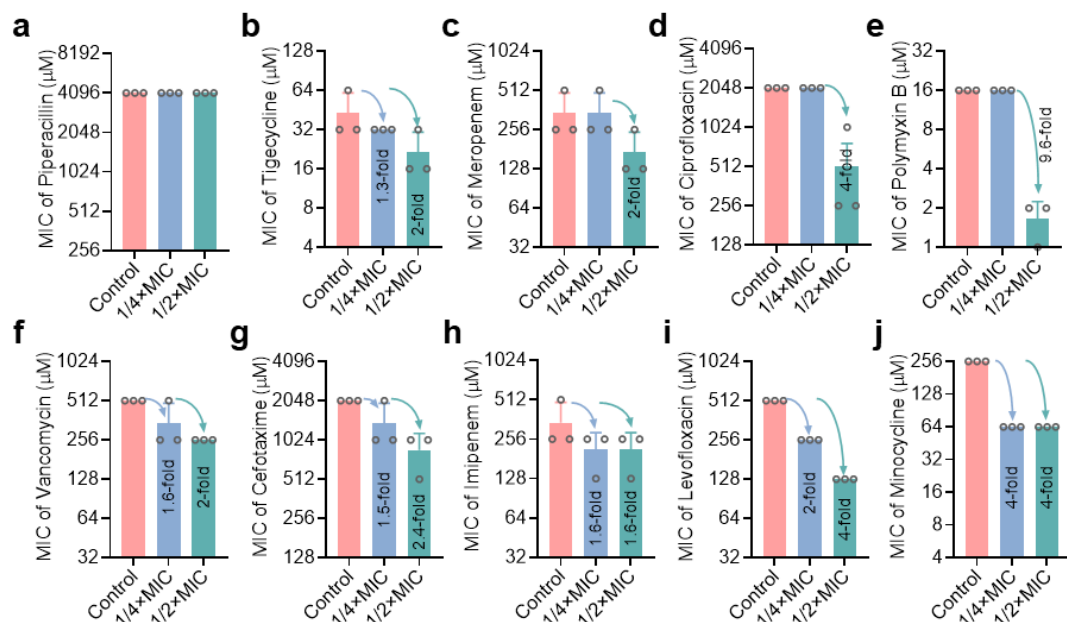
Supplementary Fig. 12 a-j Chemical sensitizations of AMP 1003 to gentamicin(a), imipenem (b), minocycline (c), tobramycin (d), levofloxacin (e), piperacillin (f), polymyxin B (g), meropenem (h), ceftazidime (i), or ciprofloxacin (j) against *MRSA* ATCC 33591, n = 3 independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.



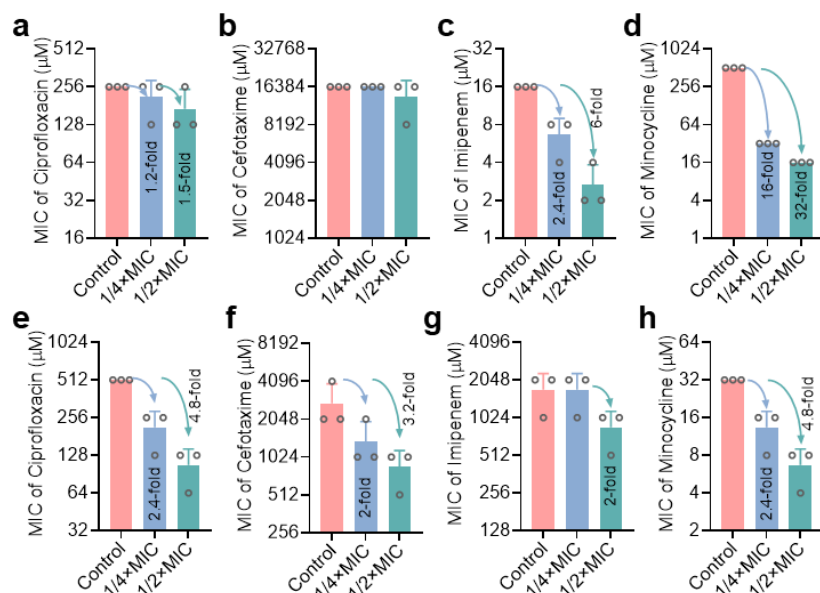
Supplementary Fig. 13 a-j Chemical sensitizations of AMP 1003 to piperacillin (a), gentamicin (b), imipenem (c), minocycline (d), ceftazidime (e), tobramycin (f), ciprofloxacin (g), levofloxacin (h), meropenem (i), or polymyxin B (j) against *MRSA* MDR2, n = 3 independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.



Supplementary Fig. 14 a-e Chemical sensitizations of AMP 1003 to ceftazidime (a), gentamicin (b), tobramycin (c), vancomycin (d) or piperacillin (e) against *K. pneumoniae* ATCC 700603, n = 3 independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.

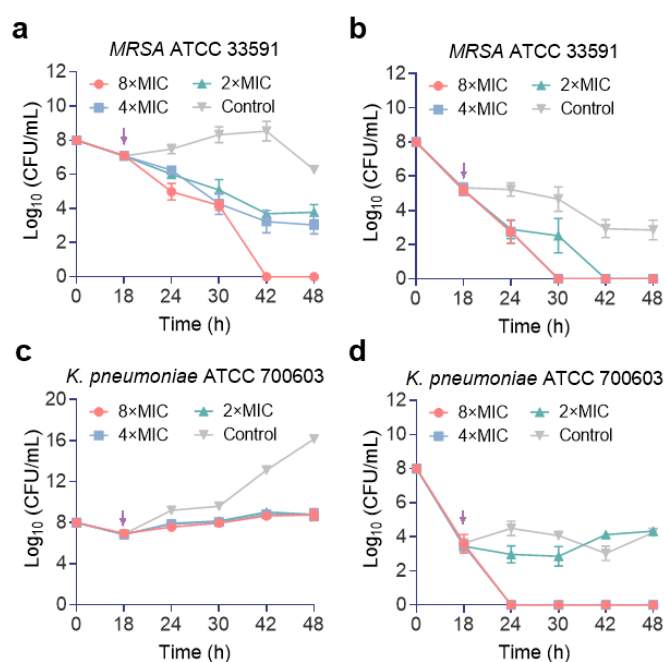


Supplementary Fig. 15 a-j Chemical sensitizations of AMP 1003 to piperacillin (a), tigecycline (b), meropenem (c), ciprofloxacin (d), polymyxin B (e), vancomycin (f), ceftazidime (g), imipenem (h), levofloxacin (i), or minocycline (j) against *K. pneumoniae* MDR5, n = 3 independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.

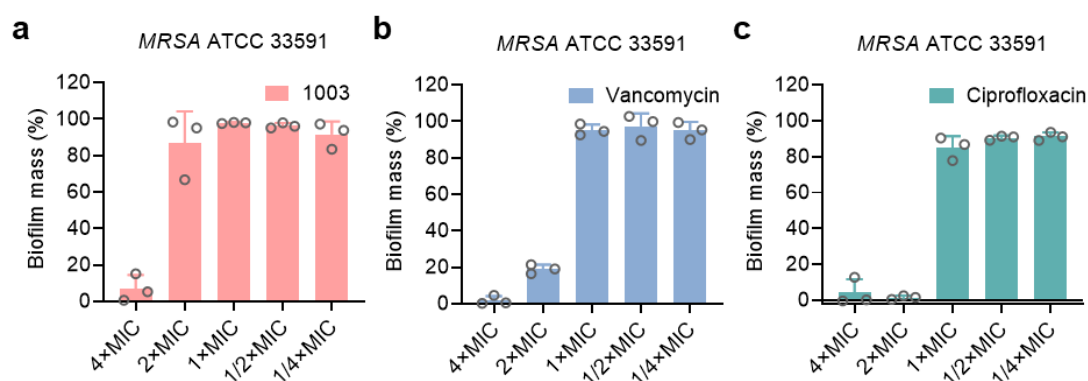


Supplementary Fig. 16 a-h Chemical sensitizations of AMP 1003 to ciprofloxacin (a), ceftazidime (b), imipenem (c), or minocycline (d) against the obtained resistant *MRSA* ATCC 33591; Chemical sensitizations of AMP 1003 to ciprofloxacin (e), ceftazidime (f), imipenem (g), or minocycline (h) against the obtained resistant *K. pneumoniae* ATCC 700603, n = 3

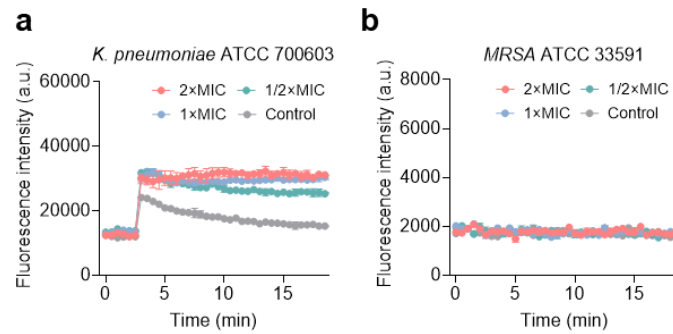
independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.



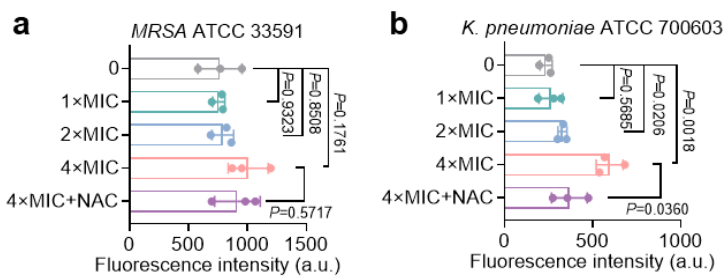
Supplementary Fig. 17 a-d Killing kinetics of AMP 1003 against *MRSA* ATCC 33591 persister cells escaped treatment with $10 \times \text{MIC}$ of ceftazidime (a) or vancomycin (b); Killing kinetics of AMP 1003 against *K. pneumoniae* ATCC 700603 persister cells escaped treatment with $10 \times \text{MIC}$ concentrations of ceftazidime (c) or polymyxin B (d), $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the median, and the error bars indicate the range.



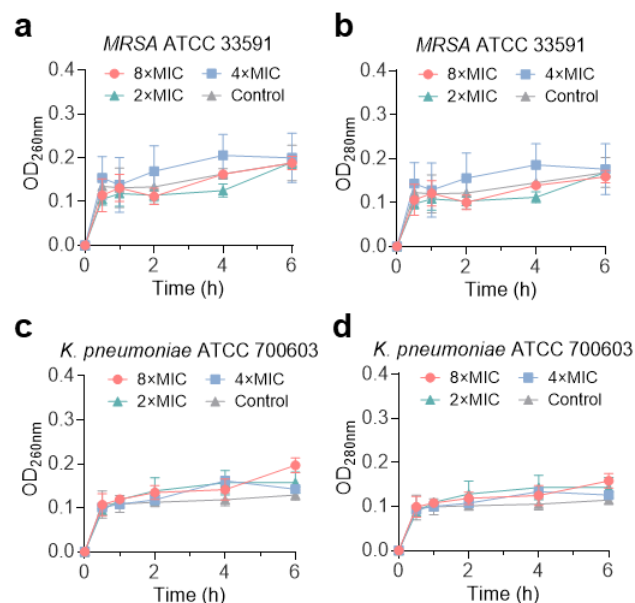
Supplementary Fig. 18 a-b The biofilm inhibition activities of AMP 1003 (a), vancomycin (b), and ciprofloxacin (c) against *MRSA* ATCC 33591, $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range.



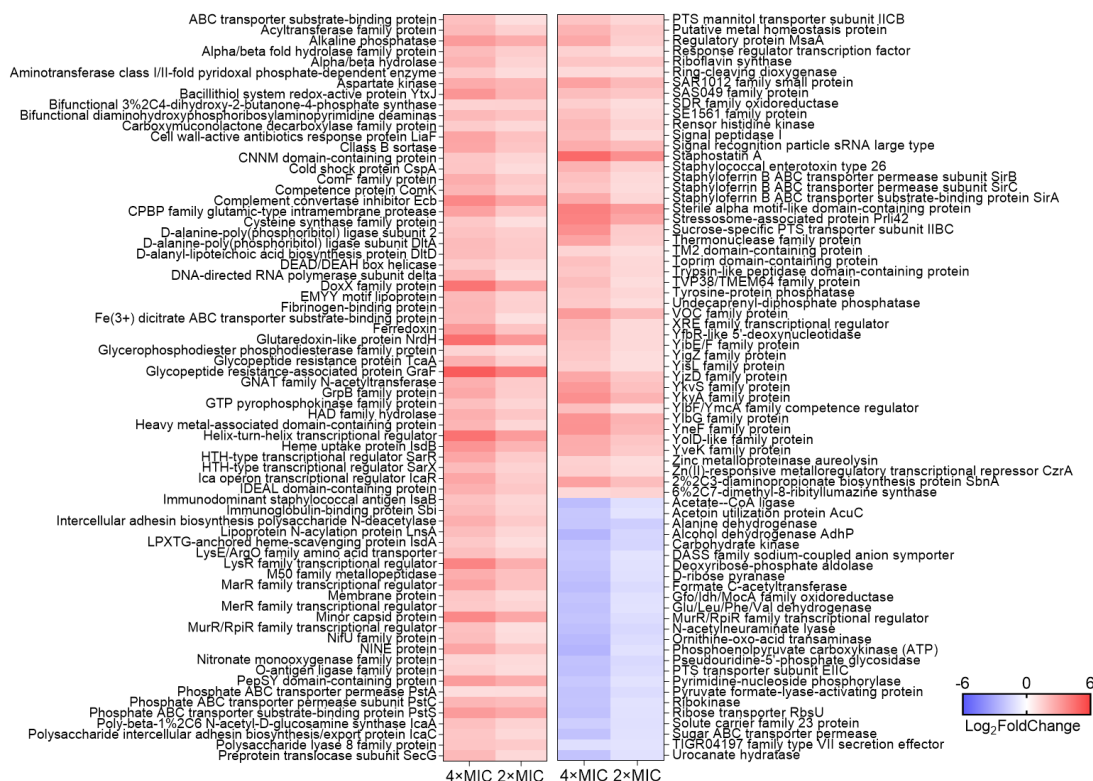
Supplementary Fig. 19 **a** Cytoplasmic membrane electrical potential of *K. pneumoniae* ATCC 700603 treated with polymyxin B at $2 \times \text{MIC}$, $1 \times \text{MIC}$ and $1/2 \times \text{MIC}$ concentrations. **b** Cytoplasmic membrane electrical potential of *MRSA* ATCC 33591 treated with vancomycin at $2 \times \text{MIC}$, $1 \times \text{MIC}$ and $1/2 \times \text{MIC}$ concentrations, $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the median, and the error bars indicate the range.



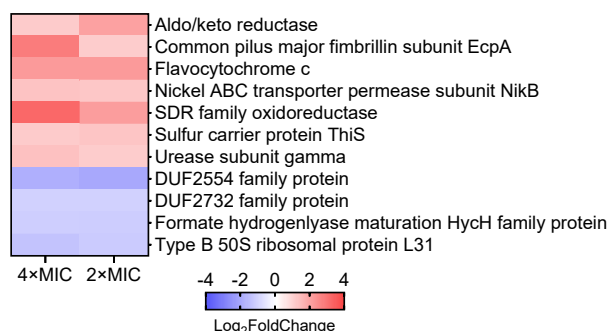
Supplementary Fig. 20 **a** ROS accumulation of *MRSA* ATCC 33591 treated with vancomycin. **b** ROS accumulation of *K. pneumoniae* ATCC 700603 treated with polymyxin B, $n = 3$ independent replicates. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. Statistical analysis: two-tailed t -test.



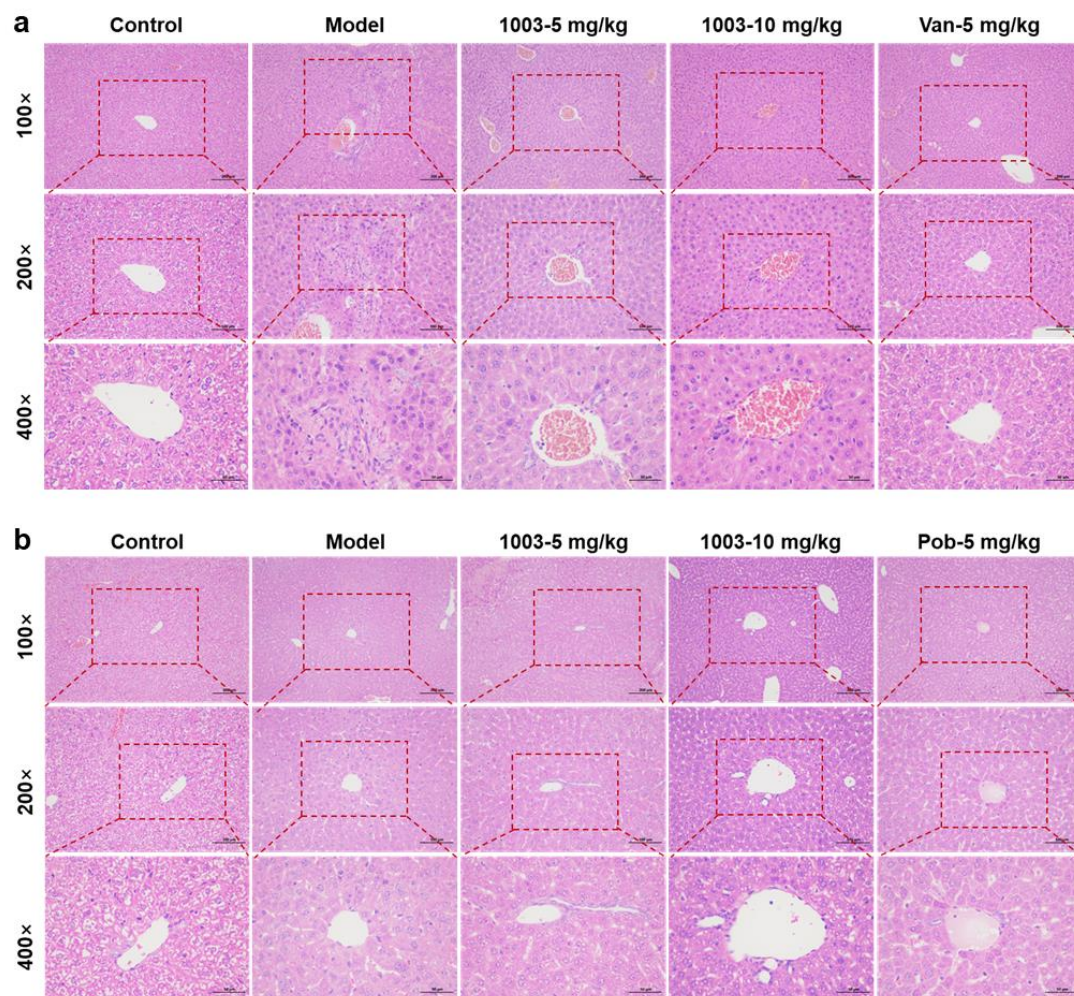
Supplementary Fig. 21 a and b Nucleic acid and protein leak of *MRSA* ATCC 33591 treated with vancomycin. c and d Nucleic acids and proteins leak from *K. pneumoniae* ATCC 700603 treated with polymyxin B, n = 3 independent replicates. Data are presented as mean values $\pm$ SD. Data points show the median, and the error bars indicate the range.



Supplementary Fig. 22 Differently expressed genes of *MRSA* ATCC 33591 under treatment with AMP 1003 at 4x MIC and 2x MIC concentrations vs. control. n = 3 biological replicates. Data are presented as means.



Supplementary Fig. 23 Differently expressed genes of *K. pneumoniae* ATCC 700603 under treatment with AMP 1003 at 4x MIC and 2x MIC concentrations vs. control. n = 3 biological replicates. Data are presented as means.

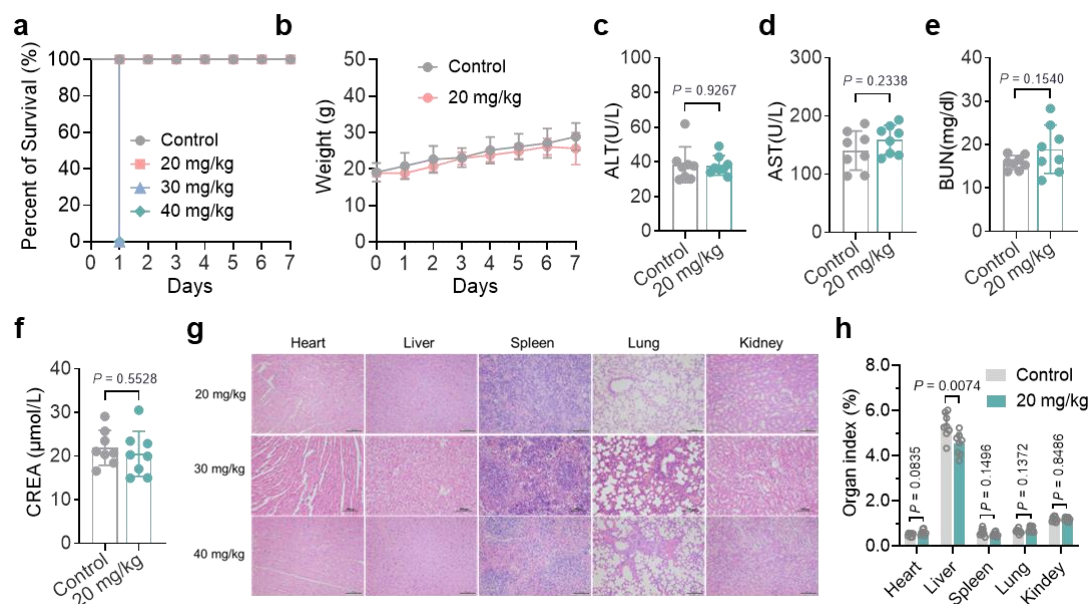


Supplementary Fig. 24 a and b Pathological sections stained by H&E of the liver in the *MRS4* ATCC 33591-infected (a) or *K. pneumoniae* ATCC 700603-infected (a) pneumonia model treated with saline, AMP 1003 (5 mg/kg or 10 mg/kg), Van (5 mg/kg) or Pob (5 mg/kg). Van stands for vancomycin. Pob stands for polymyxin B. n = 3. Scale bar: 100×: 200 μ m, 200×: 100 μ m, 400×: 50 μ m.

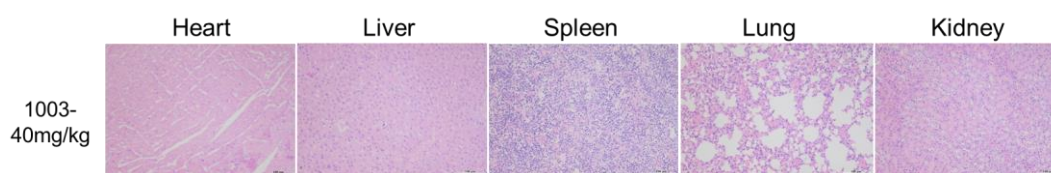
1003-20 mg/kg							Polymyxin B-20 mg/kg						
1 day	2 day	3 day	4 day	5 day	6 day	7day	1 day	2 day	3 day	4 day	5 day	6 day	7day
●	●	●	●	●	●	●	●	●	●	●	●	●	●
●	●	●	●	●	●	●	●	●	●	●	●	●	●
●	●	●	●	●	●	●	●	●	●	●	●	●	●
1003-30 mg/kg							Polymyxin B-30 mg/kg						
1 day	2 day	3 day	4 day	5 day	6 day	7day	1 day	2 day	3 day	4 day	5 day	6 day	7day
●	●	●	●	●	●	●	●	●	●	●	●	●	●
●	●	●	●	●	●	●	●	●	●	●	●	●	●
●	●	●	●	●	●	●	●	●	●	●	●	●	●
1003-40 mg/kg							Polymyxin B-40 mg/kg						
1 day	2 day	3 day	4 day	5 day	6 day	7day	1 day	2 day	3 day	4 day	5 day	6 day	7day
●	●	●	●	●	●	●	●	●	●	●	●	●	●
●	●	●	●	●	●	●	●	●	●	●	●	●	●
●	●	●	●	●	●	●	●	●	●	●	●	●	●

● Mice without signs; ● Mice with ruffled fur and poor motility; ● Mice with hutching, very ruffled fur and complete immobility even under stimulation; ● Mice died due to toxicity;

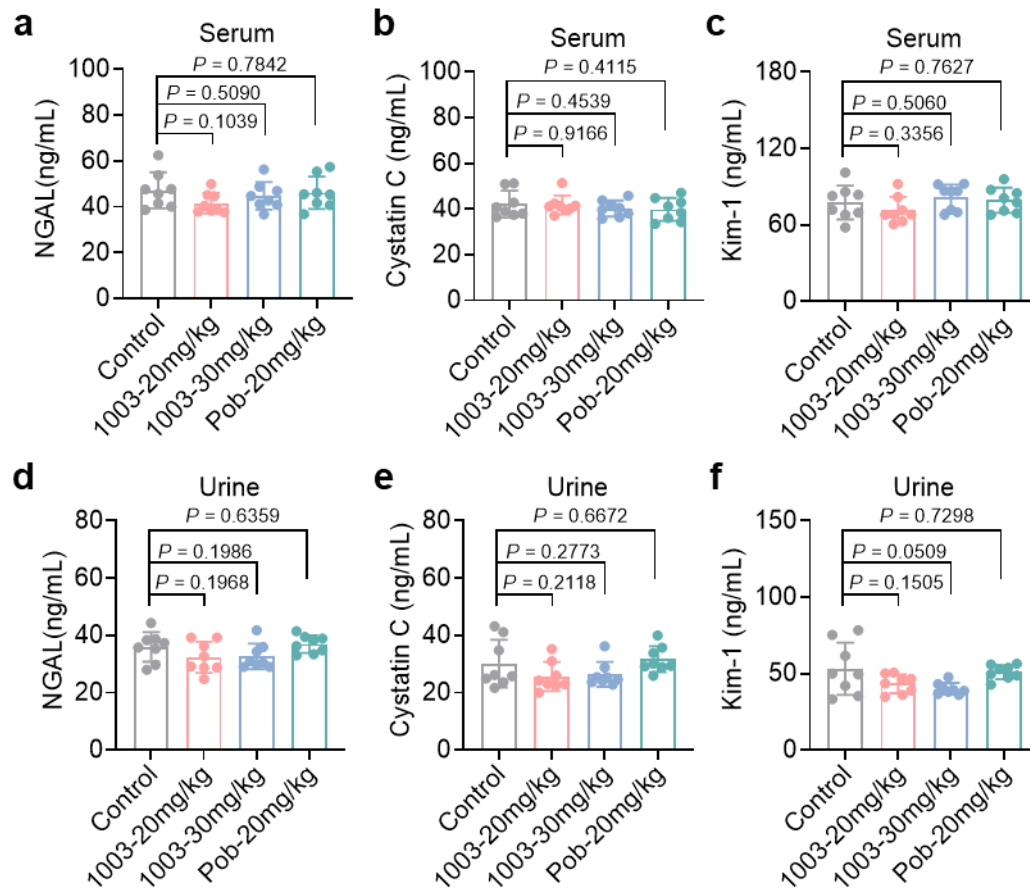
Supplementary Fig. 25 The abnormal behavior and survival of mice administered with AMP 1003 or polymyxin B for 7 days. Each circle represents one mouse, and the toxicity score was indicated by different circle shading as outlined in the legend.



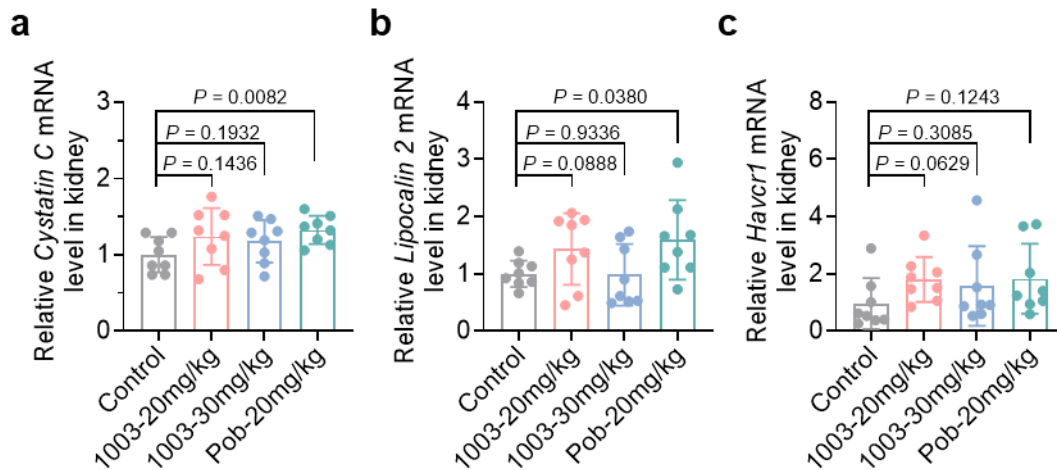
Supplementary Fig. 26 **a** Survival curves of mice treated with a single intraperitoneal injection of polymyxin B at 20 mg/kg, 30 mg/kg or 40 mg/kg. $n = 8$ mice per group. **b** The weight changes of mice administered with polymyxin B at a dose of 20 mg/kg for 7 days, $n = 8$ mice per group. Data are presented as mean values $\pm$ SD. Data points show the median, and the error bars indicate the range. **c-f** Blood biochemical analysis (ALT (**c**), AST (**d**), BUN (**e**), CREA (**f**)) of mice at 7 days post-treatment with polymyxin B at a dose of 20 mg/kg, $n = 8$ mice per group. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. Statistical analysis: two-tailed t -test. **g** Pathological sections stained by H&E of the heart, liver, spleen, lung, and kidney of the mice at 7 days after the single dose of intravenous injection with polymyxin B at a dose of 20 mg/kg, 30 mg/kg or 40 mg/kg. $n = 3$ mice per group. Scale bar: 100 μm . **h** Organ index of the mice at 7 days after the single intravenous injection dose with polymyxin B at 20 mg/kg, $n = 8$ mice per group. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. Statistical analysis: two-tailed t -test.



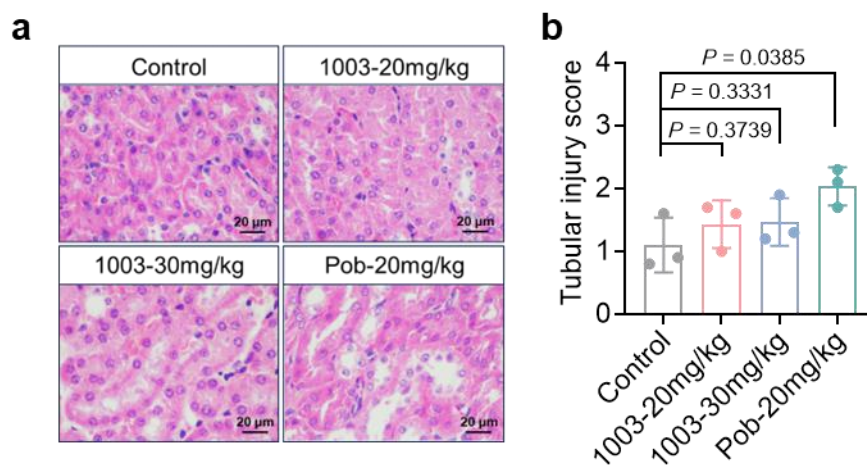
Supplementary Fig. 27 Pathological sections stained by H&E of the heart, liver, spleen, lung, and kidney of the mice died after a single dose of intravenous injection with AMP 1003 at a dose of 40 mg/kg. $n = 3$ mice per group. Scale bar: 100 μm .



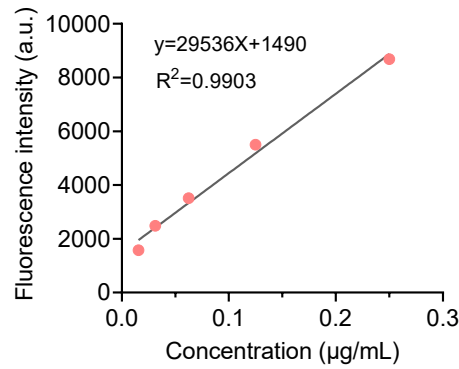
Supplementary Fig. 28 a-f The NGAL (a and d), Cystatin C (b and e) or Kim-1 (c and f) level in the serum or urine of mice at 7 days post-treatment with a single intraperitoneal injection of AMP 1003 at a dose of 20 mg/kg or 30 mg/kg and polymyxin B at 20 mg/kg, $n = 8$ mice per group. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. Statistical analysis: two-tailed t -test.



Supplementary Fig. 29 a-c The *Cystatin C* (a), *Lipocalin 2* (b), *Havcr1* (c) mRNA level in kidney of mice at 7 days post-treatment with a single intraperitoneal injection of AMP 1003 at a dose of 20 mg/kg or 30 mg/kg and polymyxin B at 20 mg/kg, $n = 8$ mice per group. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. Statistical analysis: two-tailed t -test.



Supplementary Fig. 30 a Representative (H&E) stained renal sections of mice at 7 days post-treatment with a single intraperitoneal injection of AMP 1003 at a dose of 20 mg/kg or 30 mg/kg and polymyxin B at 20 mg/kg, $n = 3$ mice per group. Scale bar: 20 μ m. **b** Mean tubular injury scores of ten randomly chosen high-power fields in kidney sections of mice at 7 days post-treatment with a single intraperitoneal injection of AMP 1003 at doses of 20 mg/kg or 30 mg/kg and polymyxin B at 20 mg/kg, $n = 3$ mice per group. Data are presented as mean values $\pm$ SD. Data points show the individual values, the central line represents the median, and the error bars indicate the range. Statistical analysis: two-tailed t -test.



Supplementary Fig. 31 Standard curve of fluorescamine.

Supplementary Tables

Supplementary Table 1: The primer for the RT-qPCR.

Gene name	Forward oligonucleotide (5'-3')	Reverse oligonucleotide (5'-3')
<i>Tnf-α</i>	CGTCAGCCGATTGCTATCT	CGGACTCCGCAAAGTCTAAG
<i>iNOS</i>	CCCTTCCGAAGTTTCTGGCAGCAGC	GGCTGTCAGAGCCTCGTGGCTTTGG
<i>Il-1β</i>	GAGAGCCGGGTGACAGTATC	TGACAAACTTCTGCCTGACG
<i>Il-17</i>	CTCCTGCTTCTAGGCTGGTTG	CCACCTGGCACTTCGAGTTAG
<i>Il-12</i>	GTGGACCAAACAATCTGACCTG	AACACGGACTATGAACCTGGA
<i>Il-18</i>	AGACTACTTCCTGAGCACAAGA	TGTCCTTACCAATGGTTCTCACT
<i>Cd86</i>	CTGGACTCTACGACTTCACAATG	AGTTGGCGATCACTGACAGTT
<i>Mcpl</i>	TTAAAAACCTGGATCGGAACCAA	GCATTAGCTTCAGATTTACGGGT
<i>Cystatin C</i>	TGGTGAGAGCTCGTAAGCAG	CATCAGATGGGGCTGGTCAT
<i>Havr1</i>	GTAAACCAGAGATTTCCACACG	TTCATGGGGACAAAATGTAGTG
<i>Lipocalin 2</i>	GACTTCCGGAGCGATCAGTT	CTGATCCAGTAGCGACAGCC
<i>β-actin</i>	CAGGATGCAGAAGGAGATTAC	AACGCAGCTCAGTAACAGTC