

## Supplementary Material

# Antibiotic synergist OM19r Reverses Aminoglycoside Resistance in Multidrug-Resistant *Escherichia coli*

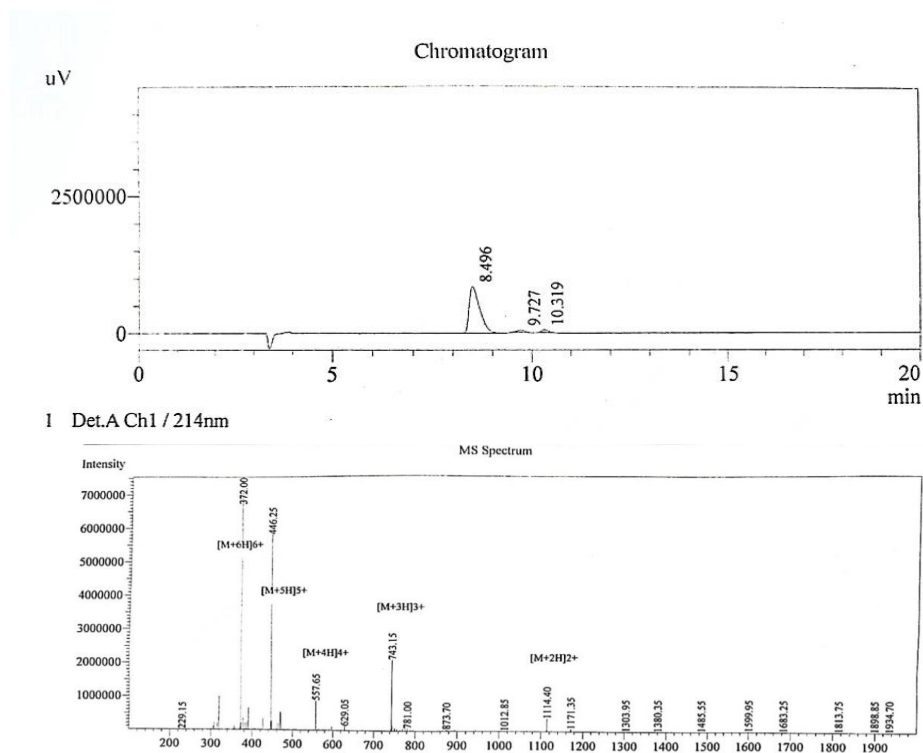
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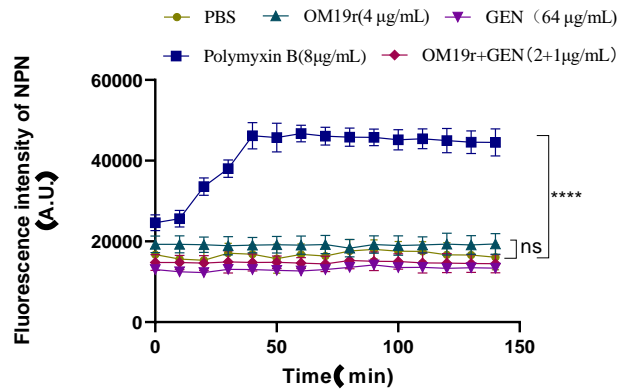
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## 1 Supplementary Figures and Tables

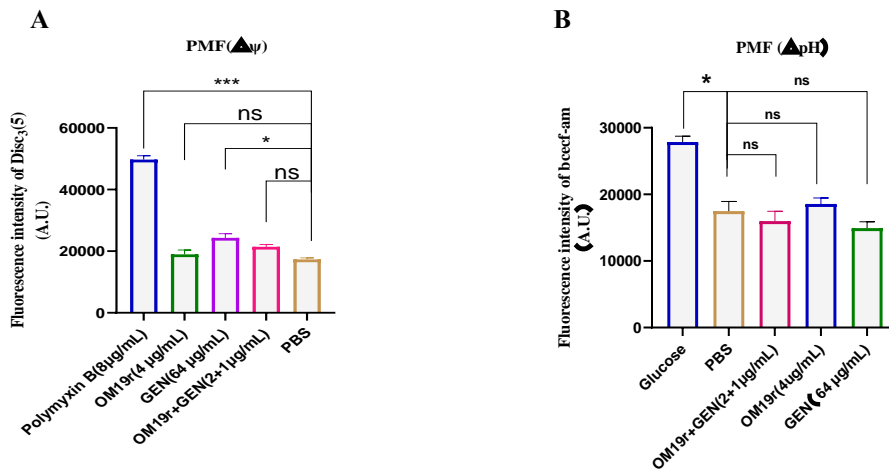
### 1.1 Supplementary Figures



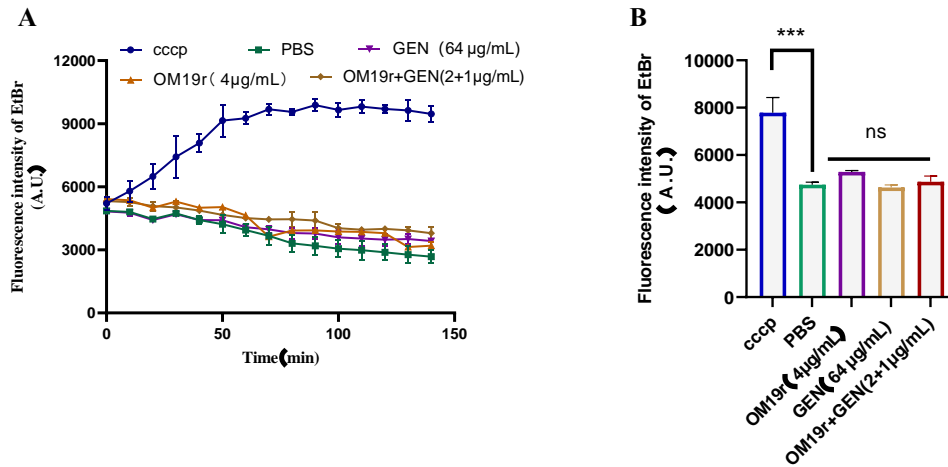
**Figure. S1** The synthetic mass spectrum of antimicrobial peptide OM19r



**Figure. S2** Outer membrane permeability dynamics probed by NPN. All experiments were performed as three biologically independent experiments and mean  $\pm$  SD are shown. P-values were determined using an unpaired two-tailed Student's t-test.



**Figure S3.** Effect of OM19r combined with GEN on PMF of *E. coli* B2. A, Dynamic changes in  $\Delta\phi$  of *E. coli* B2 were determined by 3, 3-dipropylthiodicyanide dis3 (5). B, Dynamic changes in  $\Delta\text{pH}$  of *E. coli* B2 were determined by BCECF-AM. Graphs showed the mean of three biological replicates, P-values were determined using an unpaired two-tailed Student's t-test.



**Figure S4. Effect of OM19r combined with GEN on Efflux pump of *E. coli* B2.** A, Dynamic changes in Efflux pump of *E. coli* B2 were determined by EtBr, fluorescence intensity was detected every 10 minutes for 150 minutes. B, Dynamic changes in Efflux pump of *E. coli* B2 were determined by EtBr at 30 min. Graphs showed the mean of three biological replicates, P-values were determined using an unpaired two-tailed Student's t-test.

## 1.2 Supplementary Tables

**Table S1. MIC of different antimicrobial peptides against *E. coli* ( $\mu\text{g/mL}$ )**

Antimicrobial peptide	<i>E. coli</i> ATCC25922	<i>E. coli</i> B2
OM19r	1	4
OM19R	1	4
MB7R	2	4
MB19	2	4
LR <sub>PG</sub>	2	8
LR <sub>GG</sub>	2	4
LR $\alpha$	2	4

**Table S2. FIC of different antimicrobial peptides combined with gentamicin against *E. coli* B2 ( $\mu\text{g/mL}$ )**

Antimicrobial peptide	FIC
OM19r	0.156
OM19R	0.425
MB7R	1
MB19	1
LR <sub>PG</sub>	0.5
LR <sub>GG</sub>	0.5
LR $\alpha$	0.375

**Table S3. MIC of hybrid antimicrobial peptides against *E. coli* ( $\mu\text{g/mL}$ )**

Antimicrobial peptide	Sequence	<i>E. coli</i> ATCC25922	<i>E. coli</i> B2
MDAP-2	SRDSRPVQPRVQPPPPPKQKPSIYDTPIRRPGGGRKTMYA	128	64
Oncocin	VDKPPYLPRPRPPRRIYNR-NH2	8	8
OM19R	VDKPPYLPRPRPIRRPGGR-NH2	1	4
OM19r	VDKPPYLPRPRPIRrPGGr-NH2	1	4

**Table S4. Antibacterial effect of OM19r combined with 5 antibiotics on *E. coli* B2**

Antibiotics	Action site	Combination MIC ( $\mu\text{g/mL}$ )	MIC Breakpoints ( $\mu\text{g/mL}$ )	Combined FIC
ceftriaxone	cell wall	256 (R)	4	1
enrofloxacin	DNA gyrase	128 (R)	1	1
tetracycline	30S ribosome	32 (R)	16	0.5
rifampicin	RNA polymerase	64 (R)	16	0.75
flufenicol	50S ribosome	64 (R)	32	0.75
gentamicin	30S ribosome	2 (S)	16	0.156

Note: R: resistant, S: sensitive

**Table S5. Resistance spectrum of multidrug-resistant Gram-negative bacteria selected in this study**

<b>Antibiotic(<math>\mu\text{g/mL}</math>)</b>	<b>Colistin</b>	<b>Enrofloxacin</b>	<b>Tetracycline</b>	<b>Florfenicol</b>	<b>Erythromycin</b>	<b>Kanamycin</b>	<b>Streptomycin</b>	<b>Tobramycin</b>	<b>Amikacin</b>	<b>Spectinomycin</b>	<b>Gentamicin</b>
<i>E. coil</i> ATCC 25922	0.25	1	1	2	64	1	4	1	1	1	0.25
<i>E. coil</i> B2	8	32	128	256	256	512	256	128	16	512	64
<i>S. typhimurium</i> 1A	64	32	16	8	512	256	2	8	2	256	512
<i>A. baumannii</i> JS1	0.015	256	512	128	64	64	512	64	32	512	512
<i>S. dysenteriae</i> A3	0.067	0.067	32	4	64	64	64	32	16	256	16
<i>K. Pneumoniae</i> JP20	1	128	512	256	128	>512	16	16	4	512	128

**Table S6. Antibacterial effect of OM19r combined with different aminoglycosides against drug-resistant *E. coli* B2**

<b>Antibiotic</b>	<b>MIC<sup>a</sup> (µg/mL)</b>	<b>FIC index</b>	<b>MIC<sup>b</sup> with OM19r (µg/mL)</b>	<b>Potentialiation (fold) <sup>c</sup></b>
kanamycin	512	0.375	2	256
gentamicin	64	0.157	1	64
streptomycin	256	0.188	2	128
tobramycin	64	0.375	4	64
spectinomycin	512	0.375	2	256

a/b MICs of antibiotic in the absence or presence of 0.5×MIC OM19r .

c Degree of antibiotic potentialiation in the presence of 0.5×MIC OM19r.

**Table S7. Molecular docking results**

Antimicrobial agents	Receptor	Binding Energy (kcal/mol)	Ligand RMSD (Å)
OM19r	SbmA	-262.92	204.09
OM19r	ribosome	-115.33	242.11
GEN	ribosome	-140.8	288.83