

**Dose Optimization of Colistin Combinations against Carbapenem-Resistant *Acinetobacter baumannii* from Chinese Hospital-acquired Pneumonia Patients Using *In Vitro* PK/PD Model**

Xingchen Bian<sup>1,2#</sup>, Xiaofen Liu<sup>1#</sup>, Yuancheng Chen<sup>1,3</sup>, Daijie Chen<sup>4</sup>, Jian Li<sup>5</sup>, Jing Zhang<sup>1,3\*</sup>

<sup>1</sup> Institute of Antibiotics, Huashan Hospital, Fudan University & Key Laboratory of Clinical Pharmacology of Antibiotics, National Health and Family Planning Commission & National Clinical Research Center for Aging and Medicine, Huashan Hospital, Fudan University, Shanghai, 200040, China;

<sup>2</sup> College of Life Sciences, Shanghai Normal University, Shanghai, 200234, China;

<sup>3</sup> Phase I Unit, Huashan Hospital, Fudan University, Shanghai, 200040, China;

<sup>4</sup> Shanghai Jiaotong University, Shanghai, 200240, China;

<sup>5</sup> Biomedicine Discovery Institute and Department of Microbiology, 19 Innovation Walk, Monash University, Victoria 3800, Melbourne, Australia

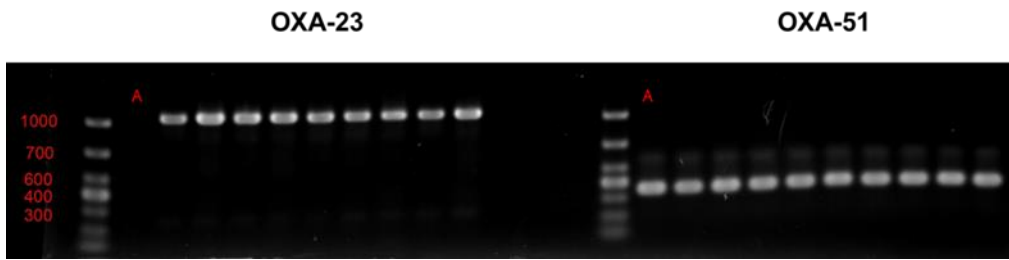
\*Corresponding author

#The authors contributed equally to the manuscript

Address correspondence to: 12 Wulumuqi Zhong Rd, Shanghai, 200040, China; Email:

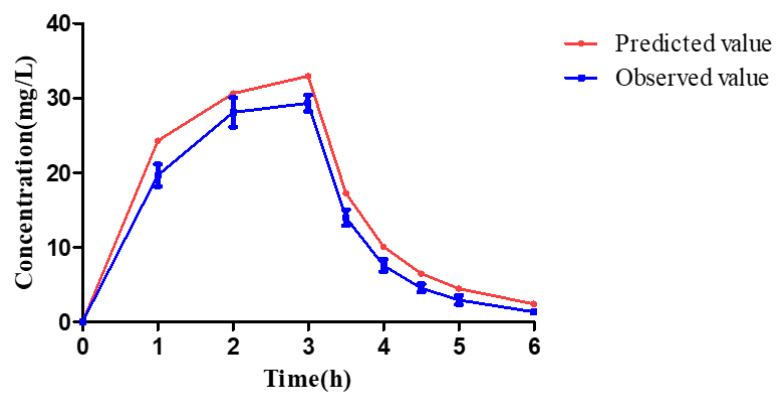
[zhangj\\_fudan@aliyun.com](mailto:zhangj_fudan@aliyun.com); Phone: +86-21-52888190.

**Key words:** *Acinetobacter baumannii*, carbapenem resistance, colistin, meropenem, combination therapy, PK/PD modeling

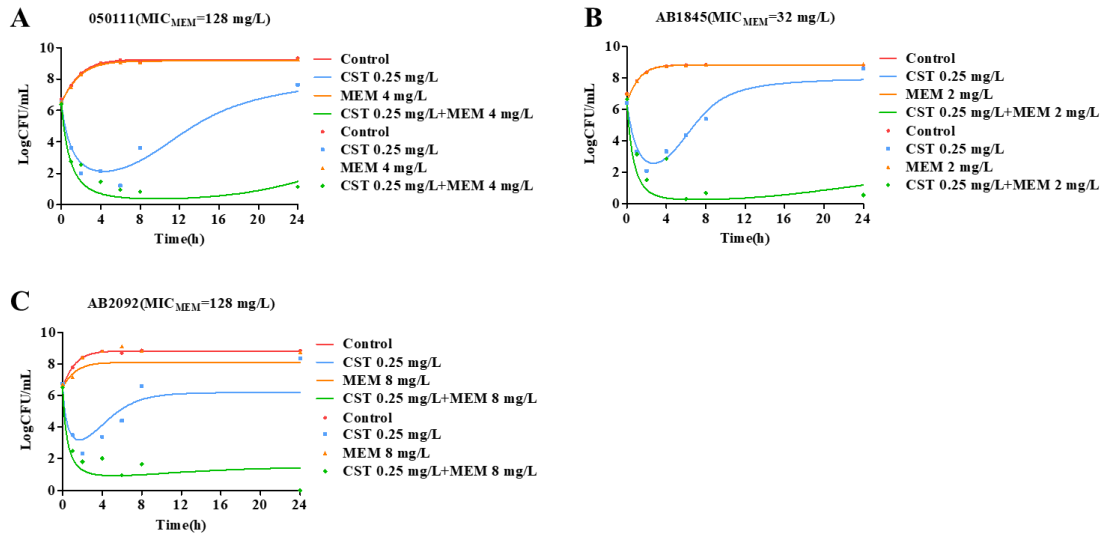


**Fig. S1** PCR results of *bla*<sub>OXA-23</sub> and *bla*<sub>OXA-51</sub> for MDR *A. baumannii* and ATCC19606

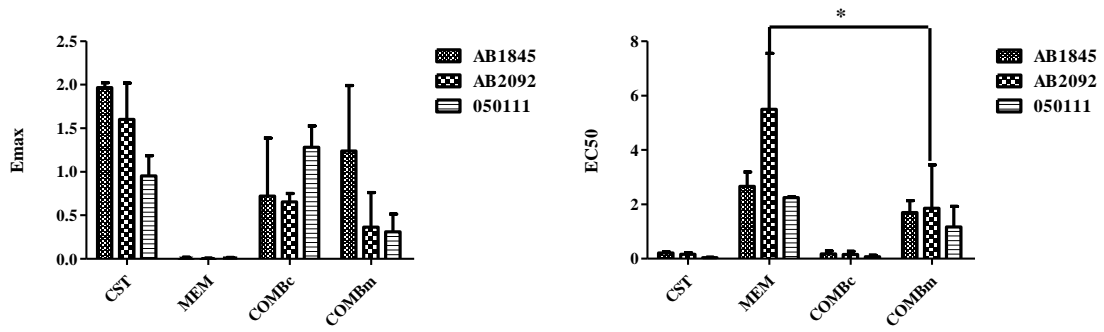
(A).



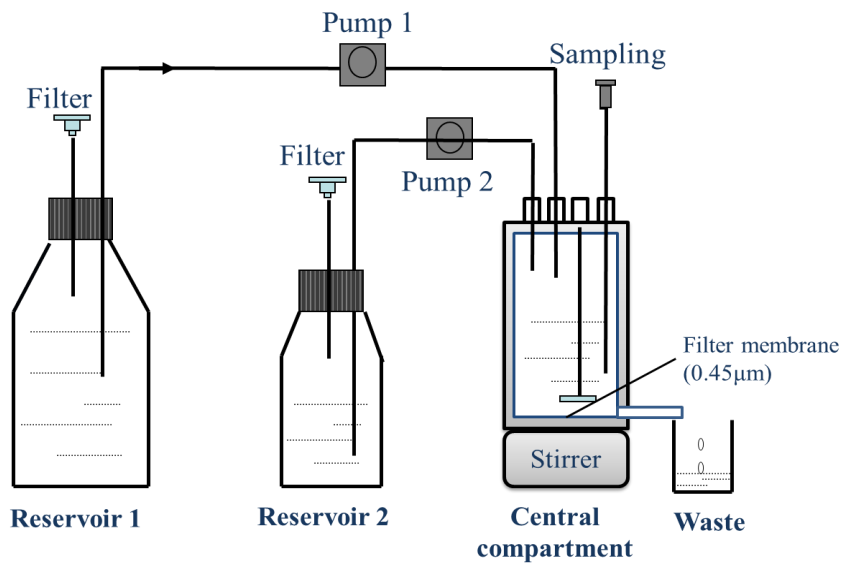
**Fig. S2** Concentration-time curves of meropenem dosed at 2 g with 3-h infusion.



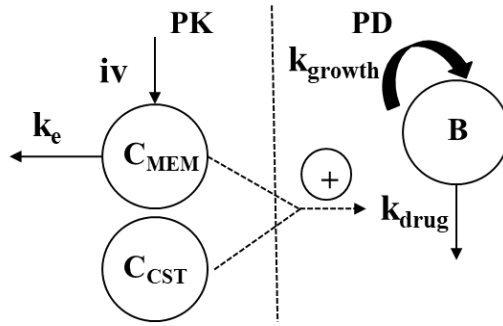
**Fig. S3** Observed (symbols) and model fitted (lines) viable counts for the static time-kill experiments with meropenem or colistin alone and in combination against *A. baumannii* 050111(A), AB1845(B) and AB2092(C). CST, colistin; MEM, meropenem.



**Fig. S4** Comparison of estimated parameters  $E_{max}$  and  $EC_{50}$  for the three strains in the static time-kill study. CST and MEM represent colistin and meropenem parameters in monotherapy; COMBc and COMBm represent colistin and meropenem parameters in the combination therapy (\*:  $p < 0.05$ ).



**Fig. S5** Schematics of the *in vitro* PK/PD model. The volume of culture medium in central compartment was 200 mL. Bacterial solution was filtered by 0.45- $\mu\text{m}$  membrane to prevent bacterial loss. The peristaltic pump was digitally controlled by a computer using WinLIN 3.2 software (Cole-Parmer Co. Ltd.). The flowing rate was regulated by section-divided modulation to mimic plasma drug concentrations in patients.



**Fig. S6** Schematics of the pharmacokinetic/pharmacodynamic (PK/PD) model characterizing the killing effect of colistin and meropenem.

**Table S1.** MICs of ATCC19606 and clinical CRAB isolates

Strain	ATCC								AB	AB
	19606	050111	020411	120211	070311	070411	080411	080511	1845	2092
CST	1	0.5	1	0.5	0.5	0.5	1	0.5	0.5	1
MEM	0.5	128	32	32	32	64	32	32	32	128
RIF	2	4	4	4	2	2	4	4	4	8
FOF	64	128	64	128	128	128	128	128	128	>128
MIN	<0.125	4	8	2	1	1	0.5	2	4	8

CST, colistin; MEM, meropenem; RIF, rifampicin; FOF, fosfomycin; MIN, minocycline; MIC, minimum inhibitory concentration.

CLSI breakpoints: CST, S:  $\leq 2$ mg/L, R:  $\geq 4$ mg/L, MEM, S:  $\leq 2$ mg/L, I: 4mg/L, R:  $\geq 8$ mg/L; MIN, S:  $\leq 4$ mg/L, I: 8mg/L, R:  $\geq 16$ mg/L.



**Table S2.** Synergistic effects of each combination against 9 clinical isolates.

<b>Combination</b>	<b>Number of strains</b>			<b>Synergy rate (%)</b>
	<b>Synergy</b>	<b>Indifference</b>	<b>Antagonism</b>	
CST-MEM	4	5	0	44
CST-RIF	5	4	0	56
CST-FOF	1	8	0	11
CST-MIN	3	6	0	33

CST, colistin; MEM, meropenem; RIF, rifampicin; FOF, fosfomycin; MIN, minocycline

Synergy rate (%) = No. of strains showing synergy / total number of strains

**Table S3.** Observed and predicted concentrations of meropenem dosed at 2 g with 3-h infusion

<b>Time(h)</b>	<b>Predicted concentration (mg/L)</b>	<b>Measured concentration (n=3) (mg/L)</b>
0	0	0.04±0.05
1	24.3	19.7±1.07
2	30.6	28.1±1.39
3	32.9	29.3±0.9
3.5	17.2	14.0±0.76
4	10.0	7.58±0.58
4.5	6.5	4.56±0.4
5	4.5	2.96±0.43
6	2.4	1.36±0.11

**Table S4.** Parameter estimates for the static time-kill study

<b>Parameter</b>	<b>Explanation</b>	<i>A. baumannii</i>	<i>A. baumannii</i>	<i>A. baumannii</i>
		<b>050111</b>	<b>1845</b>	<b>2092</b>
$k_{growth}(h^{-1})$	rate constant of bacterial net growth	0.692	0.986	0.984
$B_{max}(\log_{10}CFU/mL)$	bacterial count in the stationary phase	9.27	8.84	8.81
$E_{max\_CST}(h^{-1})$	maximum achievable kill rate constant by colistin	4.09	1.07	9.01
$EC_{50\_CST}(mg/L)$	colistin concentration that results in 50% of $E_{max}$	0.489	0.0141	0.364
$E_{max\_MEM}(h^{-1})$	maximum achievable kill rate constant by meropenem	0.014	0.719	0.16
$EC_{50\_MEM}(mg/L)$	meropenem concentration that results in 50% of $E_{max}$	4.00	7.01	2.40
$\gamma_{CST}$	hill factor for colistin	1.73	2.72	4.13
$\gamma_{MEM}$	hill factor for meropenem	30.6	5.54	$1.24 \times 10^{-7}$
$f$	maximal adaptation factor	7.82	40.7	0.561
$k$	rate of adaptation	0.0667	0.575	1.70

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<i>Int</i>	parameter describing drug interaction	5.53	9.99	2.50
<i>Fval</i> (%)	residual error fraction	1.38	3.43	2.77

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**Table S5.** Pharmacokinetic parameters of meropenem

Parameter	Explanation	Dose	
		0.5 g	1 g or 2 g
$k_e$ ( $h^{-1}$ )	elimination rate constant	1.39	1.34
$V_1$ (L)	volume of central compartment	13.7	13.8
$V_2$ (L)	volume of peripheral compartment	5.49	5.97
$k_{12}$ ( $h^{-1}$ )	speed constant from central compartment to peripheral compartment	0.506	0.323
$k_{21}$ ( $h^{-1}$ )	speed constant from peripheral compartment to central compartment	1.26	0.750