

1 **Materials and Methods**

2 **Antimicrobial agents**

3 Total of 4 antimicrobial agents were used in this study. Cefiderocol was synthesized by
4 Shionogi & Co., Ltd. (Osaka, Japan). Colistin sulfate was obtained from Wako pure
5 chemical industries, Ltd. (Osaka, Japan). Amikacin and meropenem were obtained from
6 USP (Rockville, MD).

7

8 **Test organisms**

9 Six clinical isolates (2 *P. aeruginosa*, 1 *A. baumannii*, 1 *E. coli*, and 2 *K. pneumoniae*)
10 were used in this study. All these isolates harbored at least 1 carbapenemase such as
11 VIM, IMP, OXA-23, NDM, and KPC. MIC studies were conducted with three
12 replicates, and the modal MICs were represented. For cefiderocol, iron-depleted (ID)
13 cation-adjusted Mueller Hinton broth (CAMHB) with zinc concentration of 0.65 µg/ml
14 was used as the test medium (1, 2). MICs of cefiderocol were 0.5 to 4 µg/ml. MICs of
15 colistin, amikacin, and meropenem were determined by using broth microdilution
16 method according to CLSI guidelines (2). MICs of colistin in combination with 4 µg/ml
17 amikacin were 0.125 to 2 µg/ml. All test strains showed colistin-susceptible profile with
18 colistin MIC of ≤ 2 µg/ml, however, those isolates were meropenem- and amikacin-
19 resistant profiles. MICs of meropenem and amikacin were ≥ 16 µg/ml.

20

21 ***In vitro* pharmacodynamic model**

22 We used a one-compartment *in vitro* chemostat model which is the auto-simulation
23 Shionogi dilution-type equipment (3) modified the model described by Grasso et al. (4).

24 The concentration-time curves were simulated by addition of the antimicrobial solution
25 cooled at 4°C to the bacterial suspension or by dilution of the bacterial suspension with
26 the test medium. To keep the volume of bacterial culture constant (50 ml), the culture
27 solution was automatically removed under computer control. The bacterial suspension
28 was prepared as follows. Overnight culture grown on tryptic soy agar plate was
29 suspended in saline, and then measured the optical density at 625 nm (OD₆₂₅) by
30 spectrophotometer. In this study, the number of bacterial cells was assumed to be 1×10^8
31 colony-forming unit (CFU)/ml when OD₆₂₅ was 0.1. The bacterial suspension was
32 diluted with ID-CAMHB to prepare the bacterial suspensions of 5.20×10^5 to 5.79×10^5
33 CFU/ml.

34 The antimicrobial solution was prepared as follows. Cefiderocol was dissolved and
35 diluted with saline to make 1000 µg/ml solution. Colistin and amikacin were dissolved
36 with distilled water and further diluted with saline to make 1000 µg/ml and 350 µg/ml
37 solutions, respectively. Meropenem was dissolved with 50 mM 3-(N-morpholino)
38 propanesulfonic acid buffer (pH7.0) and further diluted with saline to make 760 µg/ml
39 solution. In this study, the simulated human plasma concentrations in total 3 treatment
40 regimen (cefiderocol, colistin in combination with amikacin, and meropenem) were
41 recreated in the bacteria culture. Cefiderocol concentrations after intravenous
42 administration of 2g cefiderocol as a 3-h infusion at 8-h intervals over 24 h were
43 simulated based on the plasma concentrations after single intravenous administration of
44 2 g cefiderocol as a 1-h infusion to healthy volunteers (5), and then these concentrations
45 were corrected by the reported plasma protein binding ratio in humans (58%) (6).
46 Concentration curve of colistin was based on the 2.5 mg/kg colistin methanesulfonate

47 sodium single dose treatment administered as a 0.5-h infusion (7). The plasma
48 concentrations-time curve of amikacin is based on the 5 mg/kg single dose treatment
49 administered as a 0.5-h infusion (8). In the case of colistin, appropriate dosing regimens
50 can be designed for validation in clinical setting with robust pharmacokinetics (PK) and
51 pharmacodynamics (PD) data and associated PK/PD modeling by using the total drug
52 concentration (9). The optimal initial dosing regimen of amikacin in clinical was set by
53 PK/PD with total drug concentration (10). Meropenem concentrations after intravenous
54 administration of 1g meropenem as a 1-h infusion at 8-h intervals over 24 h were
55 simulated based on the plasma concentrations after single intravenous administration of
56 1 g meropenem to healthy volunteers (11), and then these concentrations were corrected
57 by the reported plasma protein binding ratio in humans (12.8%) (12). The PK
58 parameters to recreate the human plasma concentrations in the auto-simulation Shionogi
59 dilution- type system were calculated by compartment model analysis. Each PK
60 parameter is shown in Table S1.

61 Bacteria suspension was incubated at 37°C over 24 h under each simulated human
62 plasma concentrations-time curve. A 0.5 ml of the bacterial culture was collected at 1, 3,
63 5, 24 h for vehicle treatment and 2 h intervals over 24 h for drug treatment, diluted with
64 4.5 ml of test medium supplemented with 1 mg/ml ferric ammonium citrate, and then
65 stored at 4°C until determination of the viable cell counts. Collected samples were 10-
66 fold serially diluted with saline or ID-CAMHB. A 0.01 or 0.1 to 0.5 ml of diluted
67 bacterial cell suspensions were incubated in test medium or on agar plate (cation-
68 adjusted Mueller Hinton agar plate for meropenem and colistin/amikacin treatment or
69 ID-cation-adjusted Mueller Hinton agar for cefiderocol treatment) supplemented with 1

70 mg/ml ferric ammonium citrate, respectively, and then, the number of the colonies was
71 counted. The effect of the carry-over of the antibiotics from the culture was considered
72 to be negligible as the final concentration of the agar medium was below sub-MIC,
73 which was confirmed by the results of serially diluted samples. When the number of
74 viable cells per 1 ml of the bacterial suspension is 0, the value of \log_{10} CFU/ml
75 represents 1.

76 Table S1 Pharmacokinetic parameters of cefiderocol, colistin, amikacin, and meropenem

Drug	Human dosing regimen	Pharmacokinetic parameters			
		Vc (L)	α (h ⁻¹)	β (h ⁻¹)	kel (h ⁻¹)
Cefiderocol	2 g every 8-h as a 3-h infusion	9.846	0.763	0.268	0.518
Colistin*	2.5 mg/kg colistin methanesulfonate every 12-h as a 0.5-h infusion	1.694	0.602	0.313	0.097
Amikacin*	5 mg/kg every 8-h as a 0.5-h infusion	14.95	NA	NA	0.610
Meropenem	1 g every 8-h as a 1-h infusion	9.407	6.67	0.856	1.79

77 *: Colistin in combination with amikacin was evaluated in a one-compartment *in vitro* chemostat model.

78 Vc: distribution volume of central compartment, α : elimination rate constant in the distribution phase, β : elimination rate constant in the
 79 terminal phase, kel: elimination rate constant from central compartment, NA: not applicable.

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81 Table S2 Change of viable cells from the initial inoculum after cefiderocol, colistin/amikacin, and meropenem treatment after 24 h
 82 treatment

Strain	Initial inoculum	Change of log ₁₀ CFU/ml from the initial inoculum after 24 h treatment		
		Cefiderocol	Colistin/Amikacin	Meropenem
<i>P. aeruginosa</i>				
NUBL-7808	5.59 ± 0.08	-4.51	-2.59	+1.79
SR27001	5.20 ± 0.00	-3.30	-1.57	+2.49
<i>A. baumannii</i>				
SR08626	5.45 ± 0.10	-4.38	-2.03	+2.65
<i>E. coli</i>				
DU48916	5.79 ± 0.05	-4.53	-4.73	+2.50
<i>K. pneumoniae</i>				
SR08667	5.76 ± 0.02	-4.77	-4.73	+2.27
VA-384	5.71 ± 0.07	-4.66	-4.79	+2.67

83 CFU: colony-forming units.

84 **References**

85

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