

ORIGINAL RESEARCH

In vitro activity of colistin mono- and combination therapy against colistin-resistant Acinetobacter baumannii, mechanism of resistance, and clinical outcomes of patients infected with colistinresistant A. baumannii at a Thai university hospital

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Purpose: Colistin is a drug of last resort for treating multidrug-resistant Acinetobacter baumannii infections. Unfortunately, colistin-resistant A. baumannii (CoR-AB) has been reported. Here, we examined the in vitro effect of mono- and combined antimicrobials against CoR-AB strains and their resistance mechanism, and evaluated the clinical outcomes of CoR-AB-infected patients. Patients and methods: Seventeen clinical CoR-AB strains were isolated from patients at Phramongkutklao hospital, 2011–2015. The mono- and synergistic activities of colistin, tigecycline, sulbactam, imipenem, meropenem, amikacin, fosfomycin, and cotrimoxazole were examined by minimum inhibitory concentration (MIC) and fractional inhibitory concentration index. Clonal relationship and resistance genes were determined by repetitive extragenic palindromic polymerase chain reaction with specific primers. The effect of carbonyl cyanide 3-chlorophenylhydrazone combined with colistin was used to test efflux pump involvement. Patient treatment outcomes were also reported.

**Results:** The most prevalent infection in CoR-AB patients was pneumonia (35.3%), and all patients were administered colistin combined with another agent. The 30-day mortality was 70.6%, and the colistin MIC range and MIC50 was 16–512 μg/mL and 64 μg/mL, respectively. All CoR-AB strains were sensitive to tigecycline. Sporadic isolates were susceptible to sulbactam, imipenem, meropenem, and cotrimoxazole. A synergistic or additive effect was observed for colistin plus imipenem or meropenem (16.7%), sulbactam (66.7%), or tigecycline (66.7%). The CoR-AB isolates could be divided into four different clones (A–D) with a high prevalence of group B (47.1%). Eight isolates harbored blaOXA23, blaIMP, blaKPC, and blaNDM, and one contained blaOXA23, blaIMP, and blaKPC, while the eight remaining isolates carried only blaOXA23. The MIC values of all strains were greatly reduced for colistin plus carbonyl cyanide 3-chlorophenylhydrazone. **Conclusion:** CoR-AB clinical isolates exhibited very high colistin resistance and a high frequency of resistance genes. The mechanism of colistin resistance appears to be mediated via an efflux pump. Thus, certain antimicrobials could be used as salvage therapy for CoR-AB infection.

**Keywords:** synergism, salvage therapy, clinical outcomes

### Introduction

Acinetobacter baumannii, a gram-negative coccobacilli, is a pathogen causing major nosocomial infection, leading to a high level of unfavorable outcomes. Data from 49 US hospitals collected from 1995 through 2002 showed that A. baumannii bacteremia was

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the third highest cause of death in patients with bloodstream infections in intensive care units.<sup>1</sup> In Thailand, *A. baumannii* is the most common causative pathogen of nosocomial pneumonia in tertiary care hospitals<sup>2</sup> and the most abundant bacterium isolated from sputum cultures in a nationwide survey conducted in 55 hospitals in 2016.<sup>3</sup>

A. baumannii has been developing increasing resistance to a number of available antibiotics. Currently, the World Health Organization has defined carbapenem-resistant A. baumannii as one of the global priority superbugs. Data from National Antimicrobial Resistance Surveillance Thailand in 2016 have shown that ~70% of the clinical A. baumannii strains were resistant to carbapenems. However, no new antimicrobials are available for the treatment of carbapenem-resistant A. baumannii; thus, colistin and polymyxin B constitute the last active options in an era of multidrug-resistant A. baumannii.

Colistin was initially used during the 1950s, but was therapeutically phased out in the 1980s because of increased nephrotoxicity. However, colistin is the most effective antibiotic against multidrug-resistant gram-negative bacilli and is therefore widely used nowadays as a salvage therapy. Unfortunately, colistin-resistant *A. baumannii* (CoR-AB) emerged in 2006 as a subpopulation of the heteroresistant strains with a minimal inhibitory concentration (MIC)  $\leq 2 \mu g/mL$ . CoR-AB was first reported in the Czech republic. Data from a SENTRY study conducted across the USA, Europe, and Asia showed that CoR-AB prevalence was  $\sim 0.9\%-3.3\%$ , with levels as high as 30.6% in Korea.

The mechanism of action of colistin resistance involves the disturbance of lipid A, a component of bacterial outer membrane lipopolysaccharide. The reduced negative charge of lipid A or loss of lipid A has been hypothesized as the cause of CoR-AB. Additionally, efflux pumps, proteinaceous transporters localized in the bacterial membrane, have also been shown to significantly contribute to colistin resistance. However, to date, there are no reports of *mcr-1* in CoR-AB.<sup>6,11</sup>

To date, data regarding salvage therapy for CoR-AB infections and the clinical outcomes of patients infected with CoR-AB are limited. In addition, CoR-AB has been reported sporadically at our institute and is thus considered to be an urgent problem. Therefore, this study aimed to identify in vitro the mono- and combined antimicrobials against CoR-AB strains and their mechanism of resistance and to evaluate the clinical outcomes of patients infected with CoR-AB.

#### Patients and methods

All clinical strains and patient data were obtained from patients infected with CoR-AB admitted to Phramongkutklao

Hospital, a 1,200-bed university hospital in Bangkok, Thailand, from January 2011 to December 2015. CoR-AB was defined as *A. baumannii* with a colistin MIC >2 µg/mL as determined using the broth microdilution method according to the Clinical and Laboratory Standards Institute guidelines, 2017.<sup>12</sup>

# Ethical approval

As this study was conducted using archived bacterial isolates and involved a retrospective review of clinical outcomes, the study was given an exemption from having to obtain written informed consent from the patients by the ethical review committee of the Royal Thai Army Medical Department (approval no. Q016h/59). Following ethical approval, permission was obtained from the Director of the Phramongkutklao Hospital prior to data collection. To ensure confidentiality of patient information, anonymous typing was used and the data were maintained in a confidential manner.

# Mono- and combination antimicrobial activity

The in vitro mono-activities of tigecycline, sulbactam, imipenem, meropenem, amikacin, and cotrimoxazole were evaluated as MIC using the E-test (Liofilchem, Teramo, Italy). Colistin (Sigma-Aldrich, St Louis, MO, USA; 12 twofold dilutions [0.25–512  $\mu$ g/mL]) and fosfomycin (Wago Pure Chemical Industry, Osaka, Japan; eight concentrations [8–1,024  $\mu$ g/mL]) were evaluated using the broth microdilution and agar dilution supplemented with 25  $\mu$ g/mL of glucose-6-phosphate (Sigma-Aldrich) as the cofactor for fosfomycin transportation into the bacterial cell, respectively. *Escherichia coli* ATCC 25922 was used as the quality control strain based on Clinical and Laboratory Standards Institute guidelines. <sup>12</sup>

Checkerboard synergy testing was performed in a sterile 96-well U-bottom plate. Colistin combined with sulbactam, tigecycline, imipenem, meropenem, or cotrimoxazole (Wago Pure Chemical Industry) was examined. Each row well was inoculated with a twofold dilution of colistin (8–1,024 µg/mL), and then serial dilutions (0.25–512 µg/mL, except for tigecycline and cotrimoxazole 0.063–128 µg/mL) of the other antibacterials were added to each column well. The plates were incubated at 37°C for 18 h. MIC was defined as the lowest concentration that resulted in a clear well. Next, the fractional inhibitory concentration index ( $\Sigma$ FICI) of each combination was calculated;  $\Sigma$ FICI values  $\le$ 0.5, >0.5–1.0, >1.0–4.0, and >4.0 were interpreted as a synergistic, additive, indifferent, and antagonistic effect, respectively.

# Molecular typing and detection of resistance genes

The genomic DNA of the CoR-AB strains was extracted using a commercial kit (Thermo Fisher Scientific, Waltham, MA, USA). The clonal relationship of the CoR-AB strains was evaluated using the repetitive extragenic palindromic polymerase chain reaction (REP-PCR) method. The PCR reaction mixture was set up as previously described using the primer set REP-forward: 5′-IIIGCforwardATCAGGC-3′ and REP-reverse: 5′-ACGTCT-TATCAGGCCTAC-3′ to amplify the REP region. Thermocycling was conducted with a Biometra TGradient Thermocycler (Biometra, Gottingen, Germany). The REP-PCR products were visualized by electrophoresis on 1.2% agarose gels (Amresco, Solon, OH, USA) stained with ethidium bromide (Bio Basic, Markham, Ontario, Canada). Patterns that differed by more than three bands were classified as different clones. The commercial set of the commercial set of

The primers, PCR reaction mixture, and conditions used for the detection of carbapenemase resistance genes  $bla_{OXA-23}$  (822 bp),  $bla_{OXA-40}$  (828 bp), and  $bla_{OXA-58}$  (843 bp) were as previously described. <sup>13</sup>  $bla_{IMP}$  (232 bp),  $bla_{VIM}$  (390 bp),  $bla_{KPC}$  (798 bp), and  $bla_{NDM}$  (621 bp) were detected by multiplex PCR, as previously described. <sup>15</sup> The PCR mixture, specific primers, and PCR conditions for mcr-1 (309 bp) detection were as previously described. <sup>16</sup> All PCR products were separated by electrophoresis on 1% agarose gels (Amresco) stained with ethidium bromide (Bio Basic) and compared with the bands of known carbapenemase and mcr-1 genes (positive controls) along with the 1 kb plus DNA ladder (Thermo Fisher Scientific). PCR-grade water was used as a DNA-free negative control.

# Detection of efflux pump mechanism

Efflux pumps were evaluated using the microdilution method.  $^{12,17}$  Mueller Hinton II broth (Beckton Dickinson Company, Baltimore, MD, USA) with and without carbonyl cyanide 3-chlorophenylhydrazone (CCCP; Sigma-Aldrich) was used as the efflux pump inhibitor (EPI). CCCP was added in twofold dilutions to a colistin 96-well plate at a final concentration of 20  $\mu$ g/mL. The phenotypic determination of efflux pump activity was defined as a decreased MIC for colistin plus CCCP in more than fourfold dilution compared to the MIC of colistin alone.

## Clinical outcomes

All patients with CoR-AB infection were reviewed for clinical information including age, sex, underlying diseases, admitting ward, mechanical ventilator use, shock, hepatic function, renal function, immunocompromised status, antimicrobial

regimens, antimicrobial susceptibility of CoR-AB, source of infections, and clinical outcomes. Clinical outcomes were defined as follows: cure – clinical improvement and culture negative following treatment; clinical improvement – normal body temperature, normal level of white blood cells, and stable vital signs without microbiologic confirmation of a cure; treatment failure – worsening clinical symptoms or the requirement for different or additional antimicrobial therapy against CoR-AB; and death – in-hospital death following CoR-AB infection.

### Results

# Antimicrobial susceptibility

Seventeen cases infected with CoR-AB were identified over a 5-year period. The colistin MIC range and MIC50 of the 17 strains were 16–512  $\mu$ g/mL and 64  $\mu$ g/mL, respectively. The MIC distribution of sulbactam, tigecycline, imipenem, meropenem, trimethoprim–sulfamethoxazole, amikacin, and fosfomycin of the CoR-AB strains is detailed in Table 1.

A synergistic or additive effect was observed for colistin and imipenem or meropenem in 1/6 isolates (16.7%) and for sulbactam and tigecycline in 4/6 isolates (66.7%). No antagonistic effects were observed for any of the antimicrobial agent combinations.

# Molecular typing, resistance gene(s), and efflux pump detection

REP-PCR profiles revealed that the CoR-AB isolates could be classified into four clones, A–D. Most strains (n=8) belonged to clone B, followed by clone C (n=7), clone A (n=1), and clone D (n=1), as shown in Table 2.

Carbapenemase genes were detected by PCR; all 17 CoR-AB isolates carried at least one gene (Table 2). Importantly, a number of CoR-AB isolates harbored more than one carbapenemase gene: eight isolates contained  $bla_{OXA23}$ ,  $bla_{IMP}$ , and  $bla_{KPC}$ , and  $bla_{NDM}$ , and one contained  $bla_{OXA23}$ ,  $bla_{IMP}$ , and  $bla_{KPC}$ , while the eight remaining isolates carried only  $bla_{OXA23}$ . Interestingly, all of the isolates harboring  $bla_{OXA23}$ ,  $bla_{IMP}$ ,  $bla_{KPC}$ , and  $bla_{NDM}$  (n=8) and  $bla_{OXA23}$ ,  $bla_{IMP}$ , and  $bla_{KPC}$  (n=1) belonged to clones B and A, respectively, whereas the isolates carrying  $bla_{OXA23}$  belonged to clones C and D (Table 2).

The results of the efflux pump assay demonstrated that the colistin MIC values of the 17 CoR-AB isolates were greatly reduced (>32-fold) when combined with CCCP (Table 2).

#### Clinical outcomes

Of these 17 patients with CoR-AB infections, 11 patients (64.7%) were male, the median age was 77 years (interquartile

Table I Antibiotic minimum inhibitory concentration (µg/mL) in colistin-resistant Acinetobacter baumannii

CoR-AB isolate	COL	SUL	TIG	IMP	MEP	SMT	AMK	FOS
I	16	8	1.5	8	32	16	>4,096	256
2	32	12	2	16	32	8	>4,096	384
3	128	6	1.5	8	32	>32	>4,096	256
4	64	3	0.38	8	16	16	4,096	512
5	32	8	1.5	8	32	12	4,096	128
6	32	8	1.5	8	32	6	4,096	256
7	128	1.5	0.38	8	128	12	4,096	256
8	512	1.5	0.25	8	16	16	4,096	256
9	32	8	2	8	32	6	>4,096	384
10	32	128	1	8	32	0.38	4,096	256
H	128	32	2	32	64	>32	4,096	256
12	128	128	2	32	128	>32	>4,096	512
13	64	8	0.5	16	16	6	16	512
14	128	128	2	16	64	>32	4,096	512
15	>512	16	0.5	8	32	4	16	512
16	>512	>1,024	1	8	32	4	16	256
17	512	>1,024	2	16	128	>32	>4,096	128

Abbreviations: AMK, amikacin; COL, colistin; CoR-AB, colistin-resistant Acinetobacter baumannii; FOS, fosfomycin; IMP, imipenem; MEP, meropenem; SMT, trimethoprim-sulfamethoxazole; SUL, sulbactam; TIG, tigecycline.

**Table 2** Clonal relationship, presence of carbapenemase genes and *mcr-1*, and efflux pump detection in colistin-resistant *Acinetobacter baumannii* 

REP-PCR group	CoR-AB isolate	COL (MIC; μg/mL)	COL+CCCP	Resista	nt genes							
			(MIC; µg/mL)	bla <sub>IMP</sub>	bla <sub>NDM</sub>	bla <sub>KPC</sub>	bla <sub>OXA-23</sub>	bla <sub>OXA-40</sub>	bla <sub>OXA-58</sub>	mcr-I		
A	I	16	0.5	✓	×	✓	✓	×	×	×		
В	2	32	0.25	✓	✓	✓	✓	×	×	×		
В	3	128	0.5	✓	✓	✓	✓	×	×	×		
В	4	64	0.5	✓	✓	✓	✓	×	×	×		
В	6	32	0.25	✓	✓	✓	✓	×	×	×		
В	7	128	0.5	✓	✓	✓	✓	×	×	×		
В	8	512	0.5	✓	✓	✓	✓	×	×	×		
В	9	32	0.25	✓	✓	✓	✓	×	×	×		
В	13	64	0.25	✓	✓	✓	✓	×	×	×		
С	5	32	0.25	×	×	×	✓	×	×	×		
С	10	32	0.25	×	×	×	✓	×	×	×		
С	П	128	0.5	×	×	×	✓	×	×	×		
С	12	128	0.5	×	×	×	✓	×	×	×		
С	14	128	0.5	×	×	×	✓	×	×	×		
С	15	>512	0.25	×	×	×	✓	×	×	×		
С	17	512	0.5	×	×	×	✓	×	×	×		
D	16	>512	0.25	×	×	×	✓	×	×	×		

**Notes:** ✓ Target gene present. × Target gene absent.

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**Abbreviations:** CCCP, cyanide 3-chlorophenylhydrazone; COL, colistin; CoR-AB, colistin-resistant *Acinetobacter baumannii*; MIC, minimum inhibitory concentration; REP-PCR, repetitive element palindromic polymerase chain reaction.

range [IQR]: 62–82.5), and 11 patients (64.7%) had been admitted to the intensive care unit. The three highest ranking sources of infection were pneumonia (n=6; 35.3%), septicemia of unknown origin (n=5; 29.4%), and cholangitis (n=3; 17.6%), respectively (Table 3). All CoR-AB cases received combination antibacterial treatment, which mostly contained colistin as the backbone agent. The 30-day mortality, clinical improvement, and cure rates were 70.6%, 17.6%, and 11.8%, respectively.

## **Discussion**

CoR-AB was first isolated in 1999 and is found mostly in carbapenem-resistant strains.<sup>5</sup> Data from the EARS-Net collected from European countries revealed 5% resistance to polymyxins.<sup>11</sup> However, another study conducted in Spain revealed polymyxin resistance rates as high as 40.6%.<sup>5</sup> Fortunately, in Thailand, the National Antimicrobial Resistance Surveillance Thailand data in 2016 showed that the

Table 3 Characteristics and outcomes of patients infected with colistin-resistant Acinetobacter baumannii

Patient number	Age (years)	Sex	Comorbidity	Diagnosis	ICU	Septic shock	Prior colistin use	Regimen	Treatment duration (days)	Clinical outcome
I	58	М	DM	VAP	Yes	Yes	No	COL IMP	14	Improvement
2	81	М	DM, ESRD with HD	VAP	Yes	Yes	Yes	SUL LVX	5	Death
3	67	М	SCLC, RF	VAP	Yes	Yes	Yes	COL FOS	14	Death
4	91	F	PAD, HT, DLP, RF	Septicemia	Yes	Yes	No	COL SUL	П	Death
5	61	M	BPH, HT, RF	UTI	No	No	No	COL SUL	10	Improvement
6	90	F	HT, RF	Bacterial peritonitis	Yes	Yes	Yes	COL IMP	3	Death
7	77	М	DM, CA	Cholangitis	No	Yes	No	COL IMP	14	Death
8	81	М	HT, COPD	HAP	No	Yes	No	COL IMP	17	Improvement
9	63	F	DM, HD	Cellulitis	Yes	Yes	Yes	COL IMP	13	Death
10	82	М	HT, RF	VAP	Yes	Yes	Yes	COL MEP	12	Death
П	21	М	AKI with CVVHDF	Septicemia	Yes	Yes	No	COL MEP	4	Death
12	55	М	ALL, AKI	Septicemia	Yes	Yes	No	COL IMP	П	Death
13	80	F	HT, DM	Septicemia	No	Yes	No	COL IMP	21	Cure
14	83	М	HT, DM	Cholangitis, septicemia	No	Yes	No	COL IMP	18	Cure
15	63	F	DM	Cholangitis, septicemia	No	Yes	No	COL FOS	10	Death
16	76	М	HT, ESRD with HD	Septicemia	Yes	Yes	No	COL IMP	11	Death
17	85	F	AKI with HD	VAP with pleural effusion	Yes	Yes	No	TIG SUL	25	Death

Abbreviations: AKI, acute kidney injury; ALL, acute lymphoblastic leukemia; BPH, benign prostatic hyperplasia; CA, cancer; COL, colistin; COPD, chronic obstructive pulmonary disease; CVVHDF, continuous veno-venous hemodiafiltration; DLP, dyslipidemia; DM, diabetes mellitus; ESRD, end-stage renal disease; FOS, fosfomycin; HAP, hospital-acquired pneumonia; HD, hemodialysis; HT, hypertension; ICU, intensive care unit; IMP, imipenem; LVX, levofloxacin; MEP, meropenem; PAD, peripheral artery disease; RF, renal failure; SCLC, small cell lung cancer; SUL, sulbactam; TIG, tigecycline; UTI, urinary tract infection; VAP, ventilator-associated pneumonia.

occurrence of CoR-AB is only 0.1%;<sup>3</sup> however, this situation remains an urgent concern.

Our clinical outcome results indicate that the mortality rate was 70.6%, which is higher than the findings of Qureshi et al who showed a mortality rate of 30%. <sup>18</sup> This high mortality might be due to the fact that our patients received less active antimicrobials, even in the form of combination therapy. In contrast, half of the CoR-AB patients in the previous study received at least two or more agents, with sulbactam as the treatment regimen against CoR-AB (sulbactam MIC $_{90}$  of 8  $\mu$ g/mL). <sup>18</sup> However, data regarding the clinical outcomes of patients infected with CoR-AB are limited, and the impact of CoR-AB on morbidity and mortality is not well understood. Thus, further

clinical studies are needed to evaluate the influence of CoR-AB infection.

Our results show that the CoR-AB strains exhibited high-level colistin resistance and that colistin as a monotherapy was ineffective; however, certain antimicrobials (e.g., tige-cycline, sulbactam, or cotrimoxazole) remained active with the MIC values obtainable by standard dosages currently in clinical use. Moreover, a synergistic or additive effect was observed for colistin combined with imipenem, meropenem, sulbactam, or tigecycline. Thus, successful CoR-AB treatment could be accomplished using the remaining active agents or in combination with colistin.<sup>19</sup>

In this study, we found  $bla_{OXA23}$  in all clinical CoR-AB isolates, in agreement with previous studies conducted in

Thailand.  $^{13,20,21}$  However, to the best of our knowledge, this is the first report on  $bla_{KPC}$  identified in A. baumannii isolates in Thailand. Moreover, 52.4% of the CoR-AB isolates carried multiple carbapenemase genes. This prevalence is similar to our recent study;  $^{22}$  however, this is the first time we have identified the quadruple combination,  $bla_{OXA23}$ ,  $bla_{IMP}$ ,  $bla_{KPC}$ , and  $bla_{NDM}$ . To date, there are no existing data as to how multiple carbapenemase genes exist in A. baumannii. We hypothesize that the resistance genes might be spread via a mobile genetic element.  $^{23,24}$  However, the presence of such genes on plasmid needs to be confirmed by whole-genome or plasmid high-throughput sequencing.

A previous in vitro study has shown that CCCP significantly decreases the MIC values of colistin-resistant strains to susceptible levels (also termed colistin activity restoration). Our results suggest that colistin resistance in our CoR-AB isolates was mediated by an efflux pump. This mechanism is similar to the findings of Park and Ko;<sup>25</sup> however, CCCP cannot be used clinically because of its intrinsic cytotoxicity. To date, no active EPI has been introduced into clinical practice; however, numerous EPIs from natural compounds and of synthetic origin have been developed. Newly developed EPIs could potentially be combined with colistin as standard treatment in our region.<sup>26</sup> Additionally, to better understand the role of efflux pumps in CoR-AB, efflux pump mRNA transcription levels need to be determined.

Several limitations should be noted for our study. First, we gathered only 17 CoR-AB isolates; however, our isolates were from a 5-year period in a very low prevalence area. Second, we studied the colistin resistance via  $bla_{mcr}$  and efflux pump, but not by  $bla_{pmrAB}$  and  $bla_{lpx}$ . However, CCCP completely restored colistin activity, indicating that the resistance mechanism of all CoR-AB strains is mediated via an efflux pump. Finally, further studies are required to investigate the benefits of other active combinations against CoR-AB, such as tigecycline–sulbactam, sulbactam–imipenem, or cotrimoxazole. This in vitro study of synergism provides important information regarding the use of salvage regimens in cases of limited available antimicrobials.

### **Conclusion**

CoR-AB clinical isolates showed a very high level of colistin resistance and had a high incidence of resistance genes. However, the mechanism of colistin resistance appears to be affected via an efflux pump. With high mortality rates, certain antimicrobials could be used as salvage treatments for CoR-AB infection.

## **Author contributions**

All authors contributed toward data analysis, drafting and revising the paper and agree to be accountable for all aspects of the work.

# **Disclosure**

The authors report no conflicts of interest in this work.

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