

# Profile of ceftolozane/tazobactam and its potential in the treatment of complicated intra-abdominal infections

Marion J Skalweit<sup>1,2</sup>

<sup>1</sup>Department of Medicine,

<sup>2</sup>Department of Biochemistry, Louis Stokes Cleveland Department of Veterans Affairs and Case Western Reserve University School of Medicine, Cleveland, OH, USA

**Abstract:** Drug-resistant pathogens have gained a foothold especially in the most vulnerable patient populations, hospitalized and immunocompromised individuals. Furthermore, extended-spectrum  $\beta$ -lactamase and carbapenemase-producing organisms are finding their way even into the community, with patients presenting to the hospital with established colonization and infection with resistant Enterobacteriaceae in particular. Recently, a novel antipseudomonal cephalosporin in combination with an established Class A  $\beta$ -lactamase inhibitor, ceftolozane/tazobactam has been approved by the FDA for use in the treatment of complicated urinary tract infections and complicated intra-abdominal infections. Ceftolozane is a uniquely potent antipseudomonal cephalosporin because of its high affinity for the penicillin-binding proteins of *Pseudomonas aeruginosa*, its low affinity for the intrinsic Class C  $\beta$ -lactamases of *P. aeruginosa*, its ability to enter *P. aeruginosa* through the outer membrane without the utilization of OprD protein, and the fact that it is not a substrate of the often upregulated MexAB/OprM efflux system of *P. aeruginosa*. The biological chemistry, pharmacokinetics/pharmacodynamics, microbiologic spectrum, and clinical trials that led to the approval of ceftolozane is reviewed. A discussion regarding its potential role in the treatment of complicated intra-abdominal infections and other infectious disease syndromes associated with drug-resistant pathogens follows.

**Keywords:** ceftolozane/tazobactam, complicated intra-abdominal infections, *Pseudomonas aeruginosa*

## Introduction

The ongoing emergence of resistance among gram-negative bacteria in community- and hospital-acquired infections is inspiring the development of a variety of new  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations for the first time in decades. Among these exciting new therapeutic options is ceftolozane/tazobactam (Zerbaxa<sup>TM</sup>), a combination that contains a novel and robust antipseudomonal cephalosporin with the established Class A  $\beta$ -lactamase inhibitor, tazobactam. Ceftolozane/tazobactam has recently been FDA approved for the treatment of complicated urinary tract infections (cUTI), including pyelonephritis caused by *Escherichia coli*, *Klebsiella pneumoniae*, *Proteus mirabilis*, and *Pseudomonas aeruginosa*. In combination with metronidazole, ceftolozane/tazobactam is also indicated in the treatment of complicated intra-abdominal infections (cIAI) caused by gram-negative bacteria (*Enterobacter cloacae*, *E. coli*, *K. pneumoniae*, *Klebsiella oxytoca*, *P. mirabilis*, *P. aeruginosa*), gram-positive bacteria (*Streptococcus anginosus*, *Streptococcus constellatus*, *Streptococcus salivarius*) and anaerobic bacteria (*Bacteroides fragilis*). The microbiology of cIAI, emerging resistance in community-acquired/hospital-acquired cIAI and in cIAI in

Correspondence: Marion J Skalweit  
Department of Medicine, Louis Stokes  
Cleveland Department of Veterans  
Affairs and Case Western Reserve  
University School of Medicine, 10701 East  
Boulevard, Cleveland, OH 44106, USA  
Tel +1 216 791 3800  
Email marion.skalweit@case.edu

the immunocompromised host, as well as the potential role and significance of ceftolozane/tazobactam in the clinic as a “carbapenem-sparing” regimen are reviewed.

## Microbiology of cIAI: current guidelines for management

The microbiology of cIAI reflects the host's typical enteric flora, and generally, when a patient presents from the community, current guidelines recommend minimal testing and selection of empiric antibiotics that cover the local resistance patterns eg, CTX-M producing *E. coli*, potentially carbapenem-resistant Enterobacteriaceae (CREs).<sup>1</sup> In hospital-associated cIAI and in cIAI in immunocompromised hosts, additional pathogenic bacteria and yeast must be considered, including *P. aeruginosa* with resistance patterns that reflect local antibiograms, CREs, extended-spectrum  $\beta$ -lactamase (ESBL) producing gram-negative bacteria, and

the so-called “derepressed” AmpC hyperproducing organisms such as *E. cloacae*.

## Resistance mechanisms in pathogens found in cIAI

A variety of resistance determinants are emerging in bacteria causing cIAI, including in community-acquired infections. These include a variety of common gram-negative bacteria as well as gram-positives and anaerobic organisms. Resistance mechanisms are detailed for  $\beta$ -lactams in Table 1 and below for other agents.

## ESBL-producing organisms in the community and hospital

CTX-M-producing *E. coli* ST131 are increasing in community-acquired infections including urinary tract infections and cIAI.<sup>2,3</sup> The two main enzymes in circulation worldwide

**Table 1**  $\beta$ -lactam resistance mechanisms in bacteria associated with cIAI

Organisms	Common $\beta$ -lactam resistance determinants	Antibiotic resistance phenotype
Community-associated Enterobacteriaceae	CTX-M Class A ESBLs KPC, SME Class A carbapenemases Metallo- $\beta$ -lactamases (eg, NDM, IMP, VIM types)	Penicillins, extended-spectrum cephalosporins Penicillins, cephalosporins, carbapenems
Hospital-associated Enterobacteriaceae	TEM, SHV Class A ESBLs as well as those in community KPC, SME Class A carbapenemases Metallo- $\beta$ -lactamases (eg, NDM, IMP, VIM types) Derepressed AmpC $\beta$ -lactamases (eg, P99 or CMY-2 in <i>Enterobacter</i> spp.); Other AmpC and extended spectrum AmpC (ESAC) $\beta$ -lactamases eg, CMY Class C; $\beta$ -lactamases	Penicillins, extended-spectrum cephalosporins  Penicillins, cephalosporins, carbapenems  Penicillins, cephalosporins including cephamycins; carbapenems in OprD downregulated bacteria Cephalosporins, including cephamycins; ESACs may be more susceptible to tazobactam
<i>Pseudomonas aeruginosa</i>	Downregulation of OprD Efflux pumps (MexAB/OprM) Chromosomal AmpC of <i>P. aeruginosa</i>  Acquired ESBLs (TEM, SHV, OXA)  Acquired carbapenemases (KPC, OXA, metallo- $\beta$ -lactamases like NDM, VIM, IMP, SPM types)	Carbapenems, some cephalosporins, penicillins Meropenem, some cephalosporins, penicillins Anti <i>P. aeruginosa</i> Penicillins, anti <i>P. aeruginosa</i> Cephalosporins except ceftolozane Anti <i>P. aeruginosa</i> Cephalosporins except ceftolozane, cefepime Anti <i>P. aeruginosa</i>
<i>Acinetobacter</i> spp.	Chromosomal Class C $\beta$ -lactamases (ADC types) Acquired ESBLs Acquired carbapenemases (OXAs) Porin deletions Rnd-type efflux mechanisms	Penicillins, cephalosporins and carbapenems Penicillins, extended-spectrum cephalosporins Carbapenems Penicillins, cephalosporins, carbapenems Imipenem, tigecycline
Enterococci ( <i>E. faecalis</i> )	Altered PBPs (PBP5) Low-affinity PBPs	Ampicillin, piperacillin Intrinsic resistance to all cephalosporins and carbapenems
<i>Staphylococcus aureus</i>	Class A $\beta$ -lactamase, PCI Altered PBPs-PBP2a	Penicillin Resistance to all $\beta$ -lactams except ceftaroline fosamil

**Abbreviations:** cIAI, complicated intra-abdominal infections; ESBLs, extended-spectrum  $\beta$ -lactamases; KPC, *Klebsiella pneumoniae* carbapenemase; CTX-M, cefotaximase-Munich; SME, *Serratia marcescens* enzyme; TEM, chromosomal  $\beta$ -lactamase of *E. coli*; SHV, sulfhydryl variant of TEM; CMY-2, cephamycinase; AmpC, class C  $\beta$ -lactamase; OprD, outer porin D; Mex AB, multidrug efflux pumps AB of pseudomonas; OprM, outer porin M; OXA, oxacillinase; SPM, Sao Paulo metallo-  $\beta$ -lactamase; ADC, *Acinetobacter* derived cephalosporinase; Rnd, resistance nodulation division; PBP, penicillin binding protein; PCI, class A  $\beta$ -lactamase of *S. aureus*.

are the Type 1 CTX-M-15  $\beta$ -lactamase and the Type 9 CTX-M-14  $\beta$ -lactamase. Enterobacteriaceae (mainly *E. coli* and *K. pneumoniae*) contain CTX-M ESBLs on plasmids, and the CTX-M resistance genes can be found in association with many other resistance determinants<sup>4</sup> including fluoroquinolone resistance (*qnr* and drug modifying genes).<sup>5</sup> The presence of CTX-M ESBLs confers resistance to advanced generation cephalosporins, and in some cases, to  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations such as amoxicillin/clavulanate, ampicillin/sulbactam, ticarcillin/clavulanate, and piperacillin/tazobactam in organisms that hyperproduce these enzymes, or in the setting of an abscess (“inoculum effect”), despite the fact that CTX-M  $\beta$ -lactamases are exquisitely sensitive to inhibition by tazobactam.<sup>6</sup> Current treatment recommendations for infections caused by ESBLs, including cIAI, include carbapenems.<sup>1,7,8</sup> The widespread use of carbapenems in countries where CTX-M ESBLs are especially prevalent, as well as poor sanitation and health infrastructure, and global travel are thought to have led to the spread of CTX-M producers. It is hypothesized that this has led to more carbapenem use and the selection of even more resistant organisms that produce carbapenemases (and references therein).<sup>2</sup>

### Carbapenem-resistant Enterobacteriaceae

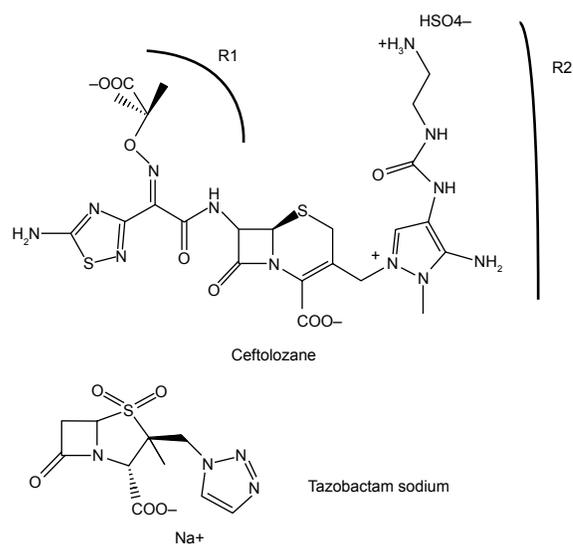
CREs are typically organisms that produce a carbapenemase such as the Class A *Klebsiella pneumoniae* carbapenemase (KPC)  $\beta$ -lactamases, Class D OXA (oxacillinase)  $\beta$ -lactamases especially OXA-48, Class B metallo- $\beta$ -lactamases eg, NDM, VIM, and IMP types. The hyperproduction of an AmpC  $\beta$ -lactamase in combination with porin loss has also been described in some CREs (reviewed by Livermore and Woodford).<sup>9</sup> These  $\beta$ -lactamases cannot be inhibited with traditional Class A  $\beta$ -lactamase inhibitors such as tazobactam. It is estimated that upward of 50% of *K. pneumoniae* isolates in Italy possess KPC carbapenem resistance determinants,<sup>10</sup> and 72% of *K. pneumoniae* isolates were imipenem resistant in Karachi, Pakistan, where KPC, OXA-48, and NDM metallo- $\beta$ -lactamases are prevalent.<sup>11</sup> Recent CRE hospital outbreaks associated with intensive care units (ICUs) and with endoscopy equipment in the US have garnered more widespread attention<sup>12–14</sup> to the problem of CREs and their role in infections. Carbapenemase genes can be found in conjunction with resistance determinants to fluoroquinolones, tetracyclines, sulfonamides, etc. Infections with CREs are associated with high morbidity and mortality,<sup>15</sup> and hospital outbreaks with such organisms are growing in frequency.<sup>16–18</sup>

### Nonfermenters, notably *P. aeruginosa*

Antibiotic resistance among *P. aeruginosa* strains is due to a variety of resistance mechanisms.  $\beta$ -lactam resistance is typically caused by the presence of upregulated efflux pumps, downregulation of OprD porin, and by the presence of specific  $\beta$ -lactamases (AmpC of *P. aeruginosa*, OXA, KPC, and metallo- $\beta$ -lactamases, Class A ESBLs). In one study of 414 ICU patients, 43% had intestinal colonization with *P. aeruginosa* at admission.<sup>19</sup> Resistance to three or more drugs was found in approximately 50% of *P. aeruginosa* isolates from the intestinal reservoir of these ICU patients who had been hospitalized 10 days or longer and was associated with disease severity and the prior use of antibiotics (fluoroquinolones, meropenem, and ertapenem).

### Ceftolozane/tazobactam

Ceftolozane (Figure 1) (formerly known as CXA-101 and FR264205) is a unique parenteral 3'-aminopyrazolium cephalosporin first discovered by Fujisawa Pharmaceutical Company in 2004.<sup>20,21</sup> It has a unique medicinal chemistry reviewed by Zhanel et al.<sup>22</sup> Ceftolozane contains the R1 side-chain of ceftazidime, another antipseudomonal cephalosporin as well as a unique R2 side-chain that confers resistance to hydrolysis by the AmpC  $\beta$ -lactamase of *P. aeruginosa* because of the reduced affinity for these enzymes.<sup>20</sup> The compound has a high affinity for the penicillin-binding proteins of *P. aeruginosa*, accounting for its excellent activity against this organism.<sup>23,24</sup> Ceftolozane is not affected by overexpression of MexAB/OprM efflux pumps because it is not a substrate of these pumps, or



**Figure 1** Chemical structure of ceftolozane/tazobactam.

**Table 2** MICs for ceftolozane/tazobactam against select bacterial causes of cIAI

Organisms	MIC range (mg/L)
<i>P. aeruginosa</i> with efflux pumps	0.5
<i>P. aeruginosa</i> with decreased OprD expression	0.5
<i>E. coli</i> CTX-M-15	0.25–2
<i>E. coli</i> SHV-5	0.25–2
<i>K. pneumoniae</i> CTX-M-15	0.25–64
<i>K. pneumoniae</i> KPC	>16
<i>Bacteroides fragilis</i>	<0.12–>16; MIC <sub>50</sub> =1

**Abbreviations:** MICs, minimum inhibitory concentrations; cIAI, complicated intra-abdominal infections.

deletion of OprD porins because it does not enter bacterial cells through OprD.<sup>25</sup>

When given alone, ceftolozane is active against many clinical strains of *P. aeruginosa*, as well as Enterobacteriaceae that do not express ESBLs or carbapenemases (Table 2 of typical minimum inhibitory concentrations [MICs]).<sup>6,20,25–30</sup> In combination with the Class A  $\beta$ -lactamase inhibitor, tazobactam, the spectrum broadens to include ESBL-producing organisms as well as some anaerobic bacteria, notably *B. fragilis*.<sup>6,20,25–30</sup> The anaerobic spectrum of ceftolozane/tazobactam is further expanded with the addition of metronidazole.<sup>29</sup>

According to the package insert, ceftolozane/tazobactam is administered parenterally 1.5 g (1 g/0.5 g) every 8 hours with metronidazole 500 mg every 8 hours for the treatment of cIAI. It must be renally adjusted (creatinine clearance [CrCl] 30–50 mL/min 750 mg IV every 8 hours; 15–29 mL/min 375 mg IV every 8 hours; end-stage renal disease or chronic kidney disease V on hemodialysis single loading dose of 750 mg followed by 150 mg IV every 8 hours, with dosing as close as possible to the end of the hemodialysis treatment session) and may have reduced efficacy in patients with CrCl  $\leq$  50 mL/min. Adverse reactions include nausea, diarrhea, headache, and fever; ceftolozane/tazobactam is contraindicated in patients with serious  $\beta$ -lactam hypersensitivity reactions. As with all antibiotics that have broad spectrum activity against enteric bacilli, ceftolozane/tazobactam can cause *Clostridium difficile*-related diarrhea.

## Pharmacokinetics/ pharmacodynamics (PK/PD) studies

A Phase I study was performed initially to evaluate the safety, tolerability, and PK of ceftolozane in healthy adult volunteers.<sup>31</sup> This placebo-controlled trial established that ceftolozane was safe and well tolerated in the 64 people studied, up to 2,000 mg in a single dose, and up to 3 g in

multiple doses (1,000 mg IV q8h and 1,500 mg q12h), for up to 10 days. This study demonstrated linear dose kinetics with a mean drug plasma half-life of 2.3 hours. More than 90% of the drug was renally eliminated, unchanged. This provided the initial safety and PK data for subsequent trials of efficacy. The PK and safety of the ceftolozane/tazobactam combination was studied in a Phase I trial involving six subjects receiving each component alone, and then the combination with doses as above.<sup>32</sup> The half-life (2.48–2.64 hours), total clearance (4.35–6.01 L/h), and volume of distribution (11.0–14.1 L) of ceftolozane were consistent with different dosing schedules and doses, and with the coadministration of tazobactam. No adverse events were noted in this study either. In summary, ceftolozane/tazobactam has linear PK and there is no drug accumulation when dosed every 8 hours. Ceftolozane is renally eliminated mostly via glomerular filtration and clearance is not affected by tazobactam, which is eliminated by renal tubular secretion.

## Clinical trials of ceftolozane/tazobactam plus metronidazole for cIAI

### Phase II

Ceftolozane/tazobactam (1.5 g IV q8h) plus metronidazole 500 mg IV q8h was compared to meropenem (1 g IV q8h)<sup>33</sup> in a prospective, double-blind, randomized, multicenter Phase II clinical trial involving adults with cIAI requiring surgical intervention (ClinicalTrials.gov, No NCT01147640). Patients were randomized in a 2:1 (ceftolozane/tazobactam + metronidazole, 82 patients vs meropenem, 39 patients) and ceftolozane/tazobactam + metronidazole was found to be non-inferior to meropenem with clinically evaluable cure rates of 91% (94% meropenem) and microbiologically evaluable cure rates of 89% (96% meropenem; 87% vs 95% with meropenem for cure of cIAI caused by *E. coli*). Adverse effect rates were similar between the two groups. This trial provided further support for the development of the drug and was conducted by Cubist Pharmaceuticals.

### Phase III

The ASPECT-cIAI trials<sup>34</sup> were two identical Phase III prospective, double-blind, randomized, multicenter clinical trials involving adults with cIAI requiring surgical intervention as well (ClinicalTrials.gov, Nos NCT01445665 and NCT01445678). These trials included a total of 993 patients primarily in Europe (76.9%). In these trials, either ceftolozane/tazobactam 1.5 g IV q8h plus metronidazole

500 mg IV q8h or meropenem 1 g IV q8h was administered for 4–14 days to hospitalized patients. Patients with staged repair with delayed fascial closure, poor source control, estimated glomerular filtration rate <30 mL/min, and antimicrobials for cIAI >24 hours prior to the first dose of study drug (unless empiric therapy had failed) were excluded from these studies. Patients were randomized in a 1:1 fashion, and received doses of study or comparator drug that were renally adjusted, using placebo saline infusions to preserve blinding of the patients and clinical staff. Intra-abdominal specimens were collected according to IDSA guidelines.<sup>1</sup> Endpoints were prospectively defined as clinical cure rates at a test-of-cure visit taking place 24–32 days from the beginning of therapy. The results were analyzed using intention to treat (ITT), microbiological intention to treat (MITT), per protocol clinically evaluable, and per protocol microbiologically evaluable groups. Patients with missing data were considered failures in the ITT analyses, and were excluded from the per protocol analyses. The non-inferiority margin was set at –10%. Adverse effects were also evaluated between the two treatment arms. The ceftolozane/tazobactam plus metronidazole was non-inferior to meropenem in the ITT group (83% vs 87.3% meropenem group, weighted difference –4.2%, 95% CI –8.91% to 0.54%); and in the secondary outcome, in the MITT group (94.2% vs 94.7% meropenem group, difference –1%, 95% CI –4.52 to 2.59). Among the 806 patients in the MITT evaluation, 525 samples (65.1%) grew *E. coli*, 76 (9.4%) *K. pneumoniae*, and 72 (8.9%) *P. aeruginosa*. The majority of infections in each group were polymicrobial. Resistant bacteria were found in equal numbers in each arm including 29 ESBL + Enterobacteriaceae. Among the 52 *P. aeruginosa* isolates found in the whole study, only three were found to be resistant to three or more antipseudomonal antibiotics. Notably, cure rates were higher in patients with infections caused by ESBL+ Enterobacteriaceae treated with ceftolozane/tazobactam plus metronidazole (95.8% [23/24] vs 88.5% meropenem [23/26]). When CTX-M-producing organisms were considered (the major ESBL family in community acquired Enterobacteriaceae, especially *E. coli*), 100% (13/13) of the microbiologically evaluable patients were cured with ceftolozane/tazobactam plus metronidazole compared to 72.7% (8/11) of the patients who received meropenem. Similar to the Phase II trial, adverse events, mainly nausea and diarrhea, were similar in the two arms.

Ceftolozane/tazobactam has been studied in Phase II and III clinical trials of cUTI (NCT 01345929) and is approved for use in this infectious disease syndrome; it is

currently being investigated for use in ventilated nosocomial pneumonia (ASPECT-NP NCT 02070757).

Ceftolozane/tazobactam affords clinicians with an additional agent that has good activity against *P. aeruginosa* and some ESBL-producing gram-negative bacteria. As empiric therapy for community-acquired cIAI, ceftolozane/tazobactam in combination with metronidazole offers a carbapenem-sparing alternative. Unfortunately, there is no evidence to show that limiting the use of carbapenems prevents the emergence of CREs and other multi-drug resistant organisms, as resistance determinants such as carbapenemases are typically found in genetic constructs that can be selected by a variety of broad spectrum antibiotics in frequent use among patients with cIAI and other serious infections.<sup>35</sup> In the setting of hospital-associated cIAI and potentially in immunocompromised hosts (febrile neutropenia with abdominal infection), ceftolozane/tazobactam may provide additional antipseudomonal coverage over combinations such as piperacillin/tazobactam. At this time, studies demonstrating the efficacy and safety of ceftolozane/tazobactam plus metronidazole to treat cIAI in immunocompromised hosts have not been conducted. Ceftolozane/tazobactam combined with metronidazole is unlikely to replace piperacillin/tazobactam or carbapenems as first-line therapy for cIAI, but more likely will be used in settings where local resistance trends suggest an advantage.

### **Future concerns/future combinations: what will this $\beta$ -lactam/ $\beta$ -lactamase inhibitor combination offer in the future compared to others waiting to enter the market?**

Since ceftolozane/tazobactam was approved for clinical use in December 2014, an additional  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combination has been FDA approved for similar clinical indications: ceftazidime/avibactam (AVYTAZ™; AstraZeneca). In this combination, a relatively less effective antipseudomonal cephalosporin is combined with a novel non- $\beta$ -lactam/ $\beta$ -lactamase diazabicyclooctane (DBO) inhibitor, avibactam (formerly NXL-104; Novoxel Pharma) to afford coverage against *P. aeruginosa* including KPC, OXA, AmpC, and ESBL-producing organisms, as well as KPC-producing CREs. DBO inhibitor combinations are not effective against organisms such as *P. aeruginosa* that harbor efflux and porin deletion resistance mechanisms. Additional combinations of avibactam with ceftaroline, an

anti-MRSA (methicillin resistant *Staphylococcus aureus*) cephalosporin and with aztreonam, an older monobactam agent with antipseudomonal activity as well as relative resistance to hydrolysis by metallo beta-lactamase, will be forthcoming. Merck Pharmaceuticals, which recently acquired Cubist Pharmaceuticals, the manufacturer of ceftolozane/tazobactam, has a similar inhibitor, relebactam in combination with imipenem–cilastatin. This combination consists of an antipseudomonal carbapenem that is not subject to known efflux resistance mechanisms in *P. aeruginosa*, together with a DBO  $\beta$ -lactamase inhibitor that will be effective in the setting of KPCs, OXA carbapenemases, and ESBLs. Imipenem–cilastatin/relebactam will not be effective against metallo- $\beta$ -lactamase producing or porin-deficient *P. aeruginosa* strains. It would seem that perhaps the best of all possible combinations is yet to come: ceftolozane/DBO inhibitor, perhaps combined with aztreonam to combat metallo- $\beta$ -lactamase producing organisms and metronidazole to provide additional anaerobic coverage. Treatment of infections with resistant *Acinetobacter baumannii* remains an additional challenge. For some CREs, combinations of colistin and tigecycline may represent an effective treatment regimen; for extensively drug resistant *P. aeruginosa*, colistin alone, or in some settings, IV fosfomycin may represent the only treatment option. For extensively drug resistant *A. baumannii*, sulbactam containing combinations, as well as colistin, tigecycline, and fosfomycin may provide effective therapy together with source control.<sup>36</sup> Other options may include the combination of carbapenems and colistin, tigecycline, or an aminoglycoside with appropriate dosing determined through PK<sup>37</sup> but these approaches need to be studied to yield best practice recommendations. Because of the complex co-expression of resistance determinants, the use of “carbapenem-sparing” agents like ceftolozane/tazobactam will likely have limited impact on the spread of CREs in the hospital.

## Summary

Ceftolozane/tazobactam combined with metronidazole is a safe and effective new antimicrobial combination approved for the treatment of complicated intra-abdominal and urinary tract infections. Its precise role as empiric and specific treatment for cIAI in the setting of the ever-evolving enteric microbiome remains to be defined.

## Disclosure

The author is an employee of the United States Department of Veterans Affairs (USDVA). Views expressed in this

review are solely those of the author, and do not reflect the opinions of the USDVA. The author has no other conflicts of interest to disclose.

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