Global Surveillance: Susceptibility of Ceftolozane-Tazobactam against Escherichia coli, Klebsiella pneumoniae, and *Pseudomonas aeruginosa* Isolates Collected from Bloodstream Infections in the United States from 2015–2017

INTRODUCTION

- Ceftolozane-tazobactam (C-T) is an antibacterial combination of a novel antipseudomonal cephalosporin and a well-established β-lactamase inhibitor
- C-T is approved in over 50 countries, including the United States, to treat complicated urinary tract infections, including acute pyelonephritis, and complicated intra-abdominal infections
- The Program to Assess Ceftolozane-Tazobactam Susceptibility (PACTS) monitors gram-negative (GN) isolates resistant to C-T worldwide
- In the current study, isolates were collected from patients hospitalized with bloodstream infections (BSIs) from 2015–2017 within the United States

MATERIALS AND METHODS

- A total of 3,377 prevalence-based BSI GN isolates were collected during 2015 to 2017 from 32 PACTS hospitals in the United States
- Isolates were tested for C-T susceptibility by CLSI broth microdilution method in a central monitoring laboratory (JMI Laboratories)
- Other antibiotics tested included amikacin (AMK), cefepime (FEP), ceftazidime (CAZ), colistin (COL), levofloxacin (LEV), meropenem (MEM), and piperacillintazobactam (PIP-TAZ)
- Antibiotic-resistant phenotypes analyzed (CLSI, 2018) for Escherichia coli (EC) and *Klebsiella pneumoniae* (KPN) included non-carbapenem-resistant extendedspectrum beta-lactamase (ESBL, non-CR) and multidrug-resistant (MDR)
- MDR was defined as resistant to at least 3 of the following classes (extendedspectrum cephalosporins, carbapenems, antipseudomonal penicillins + β -lactamase inhibitors, fluoroquinolones, aminoglycosides, gylcylcyclines, and polymyxins)
- Antibiotic-resistant phenotypes analyzed (CLSI, 2018) for *Pseudomonas* aeruginosa (PSA) included CAZ-nonsusceptible (CAZ-NS), FEP-NS, MEM-NS, PIP-TAZ-NS, and NS to all 4 β-lactam comparators tested: CAZ, FEP, MEM, and PIP-TAZ (BL-NS)
- CLSI (2018) C-T breakpoints for *Enterobacteriaceae* (ENT) are ≤2 mg/L susceptible (S), 4 mg/L intermediate (I), and ≥ 8 mg/L resistant (R); *P. aeruginosa* C-T breakpoints are $\leq 4 \text{ mg/L}$ for S, 8 mg/L for I, and $\geq 16 \text{ mg/L}$ for R
- EUCAST (2018) COL clinical breakpoints (≤2 mg/L for S) were used for ENT

Figure 1 Prevalence of species among gram-negative pathogens from bloodstream infections



Table 1 Ceftolozane-tazobactam MIC distributions for the most common Enterobacteriaceae species

	Number and cumulative % of isolates inhibited at MIC (mg/L) of: ^a											
Organism	≤0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32	>32
Escherichia	3	75	789	390	108	22	9	8	8	6	0	4
coli	(0.2)	(5.5)	(61.0)	(88.4)	(96.0)	(97.5)	(98.2)	(98.7)	(99.3)	(99.7)	(99.7)	(100.0)
Klebsiella	2	13	213	238	97	35	7	5	7	1	2	10
pneumoniae	(0.3)	(2.4)	(36.2)	(74.0)	(89.4)	(94.9)	(96.0)	(96.8)	(97.9)	(98.1)	(98.4)	(100.0)
Enterobacter		4	54	77	24	9	5	19	9	6	0	7
cloacae		(1.9)	(27.1)	(63.1)	(74.3)	(78.5)	(80.8)	(89.7)	(93.9)	(96.7)	(96.7)	(100.0)
Serratia marcescens			1 (0.7)	18 (12.9)	100 (81.0)	25 (98.0)	1 (98.6)	0 (98.6)	0 (98.6)	0 (98.6)	1 (99.3)	1 (100.0)

^a The intensity of shading is proportional to the number of tested isolates within each row that displays the indicated MIC value.

- (Figure ²
- observed in this study

RESULTS

• A total of 2.825 ENT and 552 non-enterics were isolated

- The 5 most common GN species were EC (1,422 isolates), KPN (630), PSA (344), Enterobacter cloacae (ECL; 214), and Serratia marcescens (SM; 147)

Prevalence of ENT and PSA in GN BSIs was 93.8%

– 97.5% of these GNs had a C-T MIC ≤4 mg/L

• Prevalence of CRE among GN BSIs was low with only 23/2,825 (<1%) isolates

Table 2 MIC distributions of ceftolozane-tazobactam and 4 β-lactams tested against *P. aeruginosa*

Antimicrobial	Number and cumulative % of isolates inhibited at MIC (mg/L) of: ^a								MIC	MIC				
agent	≤0.06	0.12	0.25	0.5	1	2	4	8	16	32	64	> p		WIIC ₉₀
Ceftolozane-		1	63	202	49	17	8	1				3	0.5	1
tazobactam		(0.3)	(18.6)	(77.3)	(91.6)	(96.5)	(98.8)	(99.1)				(100.0)	0.5	
Cefepime				7	101	115	32	46	29			14	0	16
				(2.0)	(31.4)	(64.8)	(74.1)	(87.5)	(95.9)			(100.0)	Ζ	10
Ceftazidime				5	95	136	49	16	10	16		17	2	16
				(1.5)	(29.1)	(68.6)	(82.8)	(87.5)	(90.4)	(95.1)		(100.0)	۷.	10
Meropenem	16	43	82	71	43	25	24	14	16	8		2	0.5 8	ο
	(4.7)	(17.2)	(41.0)	(61.6)	(74.1)	(81.4)	(88.4)	(92.4)	(97.1)	(99.4)		(100.0)		O
Piperacillin-				1		39	183	36	34	10	15	26	1	64
tazobactam				(0.3)		(11.6)	(64.8)	(75.3)	(85.2)	(88.1)	(92.4)	(100.0)	4	04

^a The intensity of shading is proportional to the number of tested isolates within each row that displays the indicated MIC value ^b Greater than the highest concentration tested.

- EC and KPN were the most commonly isolated organisms (42% and 18.7% of GN BSI, respectively)
- 98.2% of EC and 96% of KPN were susceptible to C-T (Table 1)
- %S was <90% for CAZ, FEP, and LEV
- MDR among EC and KPN GN BSIs was 8.2% and 7.8%, respectively
- Among PSA isolates, the MIC₉₀ value for C-T (1 mg/L) was 8- to 64-fold lower than the other β-lactam comparators (FEP, 16 mg/L; CAZ, 16 mg/L; MEM, 8 mg/L; PIP-TAZ, 64 mg/L; Table 2).
- Susceptibilities of C-T and comparators for the main species and resistant phenotypes are shown in Table 3



Figure 2 Percent susceptibility of C-T and comparator agents against all 2015-2017 BSI GN isolates by US division



- Against ENT, C-T (96.1%S) was more active than other cephalosporins (FEP, 89.5%S; CAZ, 87.1%S; and PIP-TAZ 93.6%S), and only MEM (99.9%S) and AMK (99.6%S) were more active
- For EC with an ESBL, non-CRE phenotype, C-T was more active (91.6%S) than CAZ (21.9%S), FEP (23.5%S), and PIP-TAZ (85.7%S)
- Against KPN with an ESBL, non-CRE phenotype, C-T was less active (81.8%S) but still more active than CAZ (19.5%S), FEP (29.9%S), and PIP-TAZ (66.2%S)
- Against PSA, C-T and COL were the most active agents tested (98.8%S and 99.7%S, respectively)
- C-T was more active than other betalactams tested
- Over 90% of isolates NS to CAZ, FEP, MEM, or PIP-TAZ were S to C-T
- C-T maintained activity (81.8%S) against 22 isolates resistant to all other betalactams tested in this study
- C-T showed some variation in susceptibility among ENT when stratified by US census division, ranging from 92.1%S in the West South Central division to 99.2%S in the New England and West North Central divisions (Figure 2)

CONCLUSIONS

- C-T demonstrated activity against the most prevalent contemporary GN isolates from BSIs in the United States
- C-T demonstrated activity against BSI ENT isolates (96.1%S), EC (98.2%S), KPN (96.0%S), and SM (98.6%S)
- For ENT, MEM and COL were the most active drugs
- Against ESBL non-CRE EC, C-T maintained activity (>90.0%S) • For PSA, C-T demonstrated activity (98.8%S); C-T and COL (99.7%S) were
- the most potent antibiotics tested Against PSA, C-T maintained activity against isolates NS to various beta-
- lactams, including isolates NS to all other beta-lactams tested in the study
- C-T (\geq 96%S) was the only beta-lactam that had \geq 90%S against EC, KPN, and PSA species
- The results of this study suggest that C-T may be a treatment option for serious GN BSIs

Organism organism g Enterobacteriace E. coli MDR ESBL, non-CRE K. pneumoniae MDR ESBL, non-CRE Enterobacter clo Serratia marcesc P. aeruginosa CAZ-NS FEP-NS MEM-NS **PIP-TAZ-NS** BL-NS d ^a CLSI (2018) ^b FUCÀST (Ź(

amalonaticus/farmeri (2), C. braakii (1), C. farmeri (1), C. freundii (8), C. freundii species complex (10), C. koseri (10), C. youngae (3), Cronobacter sakazakii (2), Enterobacter aerogenes (43), E. asburiae (1), E. cloacae (118), E. cloacae species complex (96), Escherichia coli (1,422), E. hermannii (2), Ewingella americana (1), gram-negative rods in the family Enterobacteriaceae (1), Hafnia alvei (2), Klebsiella oxytoca (109), K. pneumoniae (630), K. variicola (4), Kluyvera ascorbata (1), Kosakonia cowanii (1), Morganella morganii (30), Pantoea agglomerans (9), P. eucrina (1), Pluralibacter gergoviae (4), Proteus mirabilis (133), P. penneri (2), P. vulgaris group (2), Providencia stuartii (7), Raoultella ornithinolytica (1), R. planticola (2), Serratia liquefaciens (5), S. marcescens (147), unspeciated Citrobacter (2), unspeciated Pantoea (5), unspeciated Raoultella (5); some species are inherently resistant to COL. Nonsusceptible to all 4 B-lactam comparators tested: ceftazidime (CAZ), cefepime (FEP), meropenem (MEM), and piperacillin-tazobactam (PIP-TAZ)

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Table 3 Susceptibilities and MIC_{50/90} of ceftolozane-tazobactam and comparators tested against main organism groups in this study

/	n	% susceptible ^a (MIC _{50/90} in mg/L)										
oup		C-T	FEP	CAZ	MEM	PIP-TAZ	LEV	AMK	COLb			
ne ^c 2,825	2 825	96.1	89.5	87.1	99.0	93.6	79.2	99.6	86.1			
	2,025	(0.25/0.5)	(≤0.12/4)	(0.25/16)	(0.03/0.06)	(2/8)	(0.06/>4)	(2/4)	(0.12/>8)			
1 4 2 2		98.2	86.0	85.9	99.6	95.5	66.7	99.8	99.8			
	1,422	(0.12/0.5)	(≤0.12/>16)	(0.25/16)	(≤0.015/0.03)	(2/8)	(0.06/>4)	(2/4)	(0.12/0.25)			
	117	80.3	10.3	12.8	95.7	62.4	1.7	97.4	99.1			
		(0.5/8)	(>16/>16)	(32/>32)	(0.03/0.06)	(8/>64)	(>4/>4)	(4/16)	(0.12/0.25)			
E 251	251	91.6	23.5	21.9	100.0	85.7	20.3	99.2	99.2			
	231	(0.5/2)	(>16/>16)	(16/>32)	(≤0.015/0.06)	(4/64)	(>4/>4)	(4/8)	(0.12/0.25)			
	630	96.0	89.8	88.4	97.8	93.0	91.6	98.9	98.9			
	030	(0.25/1)	(≤0.12/4)	(0.12/16)	(0.03/0.03)	(2/16)	(0.06/1)	(1/2)	(0.12/0.25)			
49	10	57.1	10.2	6.1	71.4	44.9	32.7	87.8	89.4			
	43	(1/>32)	(>16/>16)	(32/>32)	(0.03/>32)	(32/>64)	(>4/>4)	(2/32)	(0.12/4)			
Ξ 77	77	81.8	29.9	19.5	96.1	66.2	58.4	96.1	94.7			
		(1/8)	(16/>16)	(32/>32)	(0.03/0.12)	(16/>64)	(1/>4)	(2/8)	(0.12/0.25)			
	21/	80.8	90.7	72.9	98.6	79.9	95.8	99.5	75.1			
	214	(0.25/8)	(≤0.12/2)	(0.25/>32)	(0.03/0.12)	(2/64)	(≤0.03/0.5)	(1/2)	(0.12/>8)			
one	1/7	98.6	98.6	98.6	98.6	95.2	98.0	100.0	0.7			
5013	147	(0.5/1)	(≤0.12/0.25)	(0.25/0.5)	(0.06/0.06)	(2/8)	(0.12/0.5)	(2/4)	(>8/>8)			
34	311	98.8	87.5	87.5	81.4	85.2	77.9	97.4	99.7			
	344	(0.5/1)	(2/16)	(2/16)	(0.5/8)	(4/64)	(0.5/>4)	(4/8)	(1/1)			
13	13	90.7	27.9		32.6	11.6	41.9	88.4	100.0			
		(2/4)	(16/>16)	(32/>32)	(8/32)	(>64/>64)	(>4/>4)	(8/32)	(1/1)			
43	13	90.7		27.9	30.2	16.3	25.6	88.4	100.0			
	40	(2/4)	(16/>16)	(32/>32)	(8/32)	(>64/>64)	(>4/>4)	(8/>32)	(1/1)			
	64	93.8	53.1	54.7		46.9	31.2	89.1	100.0			
	04	(1/4)	(8/>16)	(8/>32)	(8/32)	(32/>64)	(>4/>4)	(8/32)	(1/1)			
	51	92.2	29.4	25.5	33.3	0.0	35.3	90.2	100.0			
		(1/4)	(16/>16)	(32/>32)	(8/32)	(>64/>64)	(>4/>4)	(8/16)	(1/1)			
	22	81.8					13.6	90.9	100.0			
		(2/>32)	(16/>16)	(32/>32)	(16/32)	(>64/>64)	(>4/>4)	(8/16)	(1/1)			

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