Susceptibility Trends of Ceftolozane-Tazobactam and Comparators When Tested against US Gram-Negative Bacterial Surveillance Isolates Collected from 2012–2018

Introduction

- Ceftolozane-tazobactam (C-T) is an antipseudomonal cephalosporin combined with a β -lactamase inhibitor
- -C-T has activity against most common β -lactam resistance mechanisms employed by *Pseudomonas aeruginosa* (PSA), including AmpC production (PDC), up-regulated efflux pumps, and porin reductions (OprD loss)
- -C-T also has activity against most extended-spectrum β -lactamase (ESBL)-producing Enterobacteriaceae
- C-T has been approved in >60 countries for treatment of complicated urinary tract infections and acute pyelonephritis and for complicated intra-abdominal infections (with metronidazole) in adults
- C-T was recently approved by the FDA for the treatment of hospital-acquired bacterial pneumonia, including ventilatorassociated bacterial pneumonia
- The Program to Assess Ceftolozane-Tazobactam Susceptibility (PACTS) monitors C-T resistance worldwide
- We analyzed resistance trends in the United States over the 7 years of PACTS for gram-negative (GN) isolates collected in 35 US medical centers

Materials and Methods

- In 2012–2018, 35,514 GN isolates, including 26,707 Enterobacteriaceae (ENT) and 6,583 PSA isolates, were tested for susceptibility (S) by the CLSI broth microdilution method at JMI Laboratories
- Infection types were bloodstream (BSI), patients hospitalized with pneumonia (PHP), skin and skin structure (SSSI), complicated intraabdominal (cIAI), and complicated urinary tract (cUTI)
- Antimicrobials tested included C-T, amikacin (AMK), colistin (COL), cefepime (FEP), ceftazidime (CAZ), levofloxacin (LEV), meropenem (MEM), and piperacillin-tazobactam (PIP-TAZ)
- Phenotypes identified were:

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-Carbapenem-resistant ENT (CRE),

-ENT screen-positive for ESBL that are non-CRE (ESBL)

- -CAZ nonsusceptible, MEM nonsusceptible, FEP nonsusceptible, PIP-TAZ nonsusceptible, and nonsusceptible to all 4 β -lactam (BL) comparators
- -Multidrug-resistant (MDR) isolates were identified as nonsusceptible to 3 or more antimicrobial classes
- -Extensively drug-resistant (XDR) isolates were identified as nonsusceptible to all but 2 or fewer antimicrobial classes
- CLSI (2019) interpretive criteria were used, except for ENT with COL, which used EUCAST (2019)

Results

• The most common infection type caused by a GN pathogen was PHP (10,406) followed by UTI (8,683), BSI (7,565), SSSI (5,595), and IAI (3,000) (Figure 1)

Table 1 Overall antimicrobial activity of ceftolozane-tazobactam, amikacin, cefepime, ceftazidime, colistin, levofloxacin, meropenem, and piperacillin-tazobactam tested against the main organisms and organism groups

Organism/organism g (no. of isolates) Pseudomonas aeruginose Ceftolozane-tazobactan Amikacin (6,583) Cefepime (6,580) Ceftazidime (6,583) Colistin (6,583) Levofloxacin (6,579) Meropenem (6,575) Piperacillin-tazobactam Enterobacteriaceae Ceftolozane-tazobactam Amikacin (26,658) Cefepime (26,698)

Ceftazidime (26,705)

Colistin^b (26,542)

Levofloxacin (26,680)

Meropenem (26,705)

Piperacillin-tazobactam

^b FUCAST 2019 Green is susceptible.

Yellow is intermediate or susceptible dose dependent (cefepime

- and UTI
- -PSA was the most common pathogen in PHP and SSSI C-T and comparator MIC distributions for PSA and ENT are shown in
- Table 1
- Overall, the %S of C-T vs. 1,258 MDR PSA was 86.6% and was 81.8% against 777 XDR
- respectively
- -COL %S for MDR was 98.9% and for XDR was 98.5%
- –During the period, the MDR ENT rate varied from 8.5% to 10.1% and the XDR remained flat
- -The CRE and ESBL, non-CRE rates varied from 0.9% to 2.2% and 10.5% to 16.8%, respectively
- -The PSA MDR rate increased from 16.1% to 22.9%
- -The PSA XDR rate increased from 9.5% to 15.6%
- -The BL-NS PSA rate doubled from 5.0% in 2015 to 10.3% in 2018
- Over the period, the most active drugs were
- For PSA: C-T (96.4–98.8%S), AMK (93.8–98.0%S), and COL (98.4%–100.0%) (Figure 3)

				No.	and cumu	lative % of	isolates in	hibited at	MIC (mg/L)	of ^a					
roup															
	≤0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32	64	> a	MIC ₅₀	MIC ₉₀
osa															
am (6,583)	4	12	43	588	3,576	1,479	456	240	86	26	13		60	0.5	2
	0.1	0.2	0.9	9.8	64.2	86.6	93.5	97.2	98.5	98.9	99.1		100		
				58	180	481	2,143	2,192	965	312	102		150	4	8
				0.9	3.6	10.9	43.5	76.8	91.4	96.2	97.7		100		
					220	1,105	2,129	1,024	1,078	634			390	2	16
					3.3	20.1	52.5	68.1	84.4	94.1			100		
				52	149	1,298	2,572	979	483	283	306		461	2	32
				0.8	3.1	22.8	61.8	76.7	84	88.3	93		100		
					2,191	2,795	1,560	31	0				6	1	2
			000	4.040	33.3	75.7	99.4	99.9	99.9				100	0.5	
			268	1,048	2,333	693	608	472					1,157	0.5	>4
		ΕΛΛ	4.1	20	55.5	66 740	75.2	82.4	115				100	0.5	0
		544	844	1,399	1,274	748	438 79.8	385 85.7	415				528	0.5	8
m(6.580)		8.3	21.1	42.4	61.8 260	73.1 131	460	2,669	⁹² 1,048	668	373	264	100 707	Λ	>64
ım (6,580)					200	5.9	12.9	53.5	69.4	79.6	85.2	89.3	100	4	-04
					- -	0.0	12.0	00.0	00.4	10.0	00.2	00.0	100		
am (26,707)	19	374	7,574	10,539	4,850	1,402	509	344	322	215	167		392	0.25	1
	0.2	1.5	29.8	69.3	87.5	92.7	94.6	95.9	97.1	97.9	98.5		100	0.20	•
				51	1,119	8,670	10,920	4,429	950	248	200		71	2	4
				0.2	4.4	36.9	77.9	94.5	98.1	99	99.7		100	_	
					23,239	444	392	331	309	316			1,667	≤0.5	2
					87	88.7	90.2	91.4	92.6	93.8			100		
	573	3,198	8,625	6,828	2,504	809	428	343	387	639	770		1,601	0.25	16
	2.2	14.1	46.4	72	81.4	84.4	86	87.3	88.7	91.1	94		100		
					20,569	789	116	68	88				4,912	≤0.5	>8
					77.5	80.5	80.9	81.2	81.5				100		
			18,280	1,357	1,204	578	356	464					4,441	≤0.12	>4
			68.5	73.6	78.1	80.3	81.6	83.4					100		
		25,212	799	134	58	41	65	79	73				244	≤0.06	≤0.0
		94.4	97.4	97.9	98.1	98.3	98.5	98.8	99.1	000	400	440	100	0	
ım (26,675)					2,658	4,674	10,518	4,297	1,488	868	460	449	1,263	2	16
					10	27.5	66.9	83	88.6	91.9	93.6	95.3	100		

 The top 5 GN species from the infection types are shown in Figure 1 -Escherichia coli was the most common species among BSI, IAI,

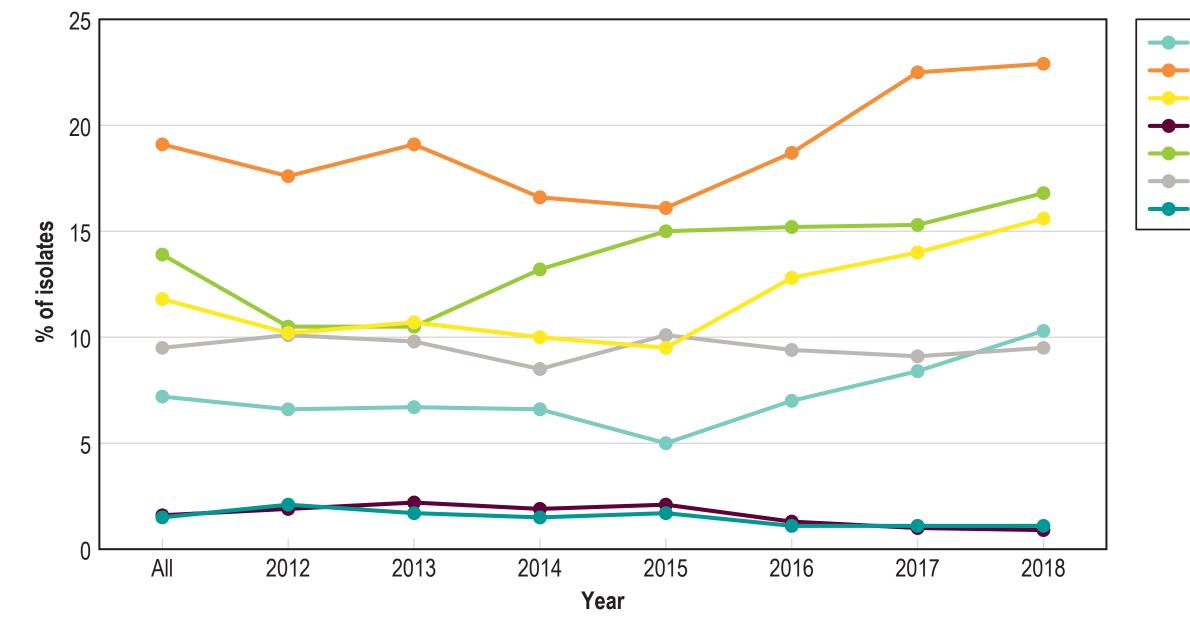
- -Against MDR and XDR PSA, AMK had 84.5% and 80.4% S,
- Frequencies of the resistant phenotypes are shown in Figure 2



Figure 1 Top 5 gram-negative isolates by infection type

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Figure 2 Change in percent of resistant phenotypes for *P*. aeruginosa and Enterobacteriaceae over the 7-year period



PSA, P. aeruginosa; MDR, multidrug-resistant; XDR, extensively drug-resistant; CRE, carbapenem-resistant Enterobacteriaceae: ESBL, extended-spectrum B-lactamase: ENT. Enterobacteriaceae

Figure 3 Susceptibility of P. aeruginosa by year

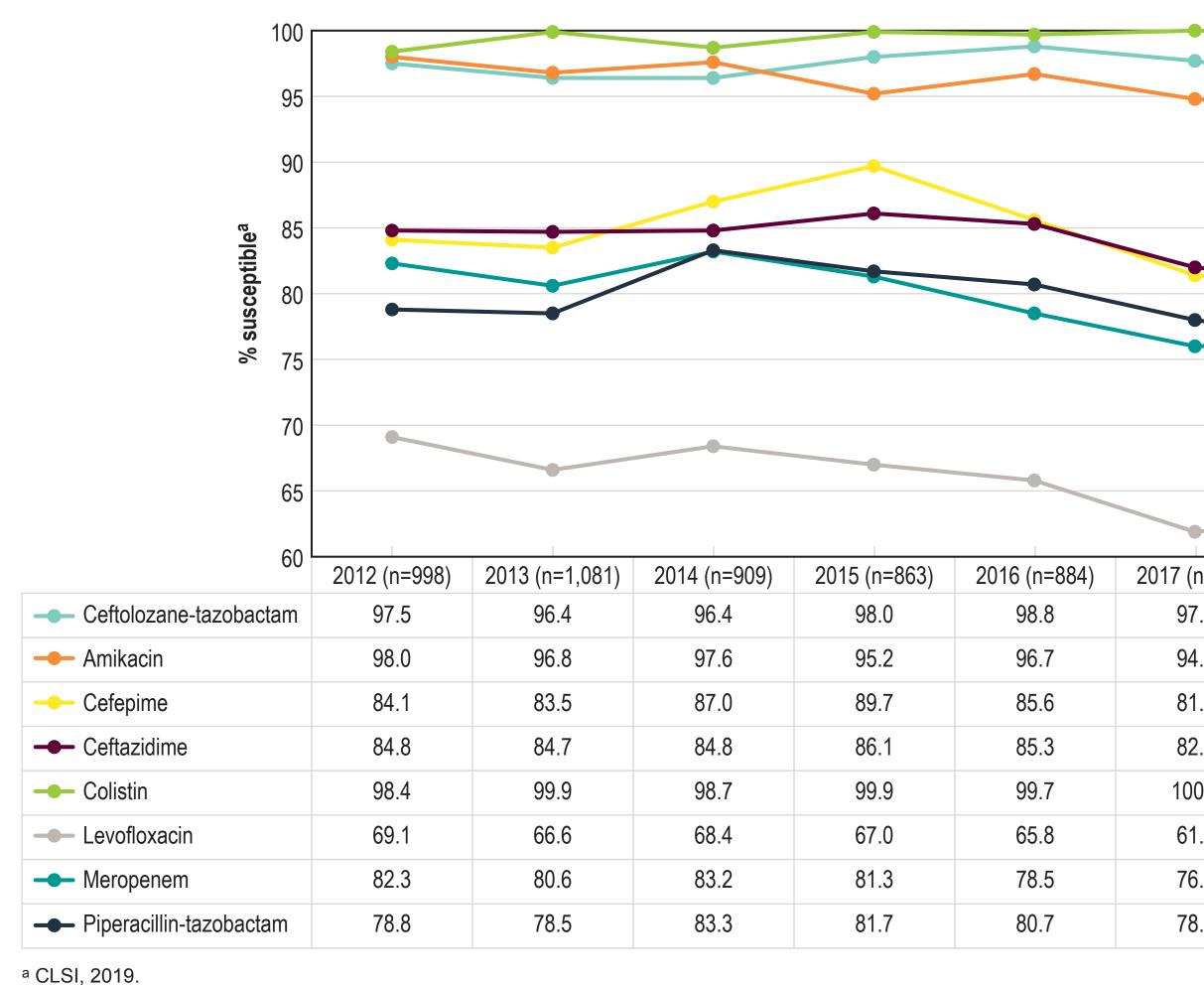
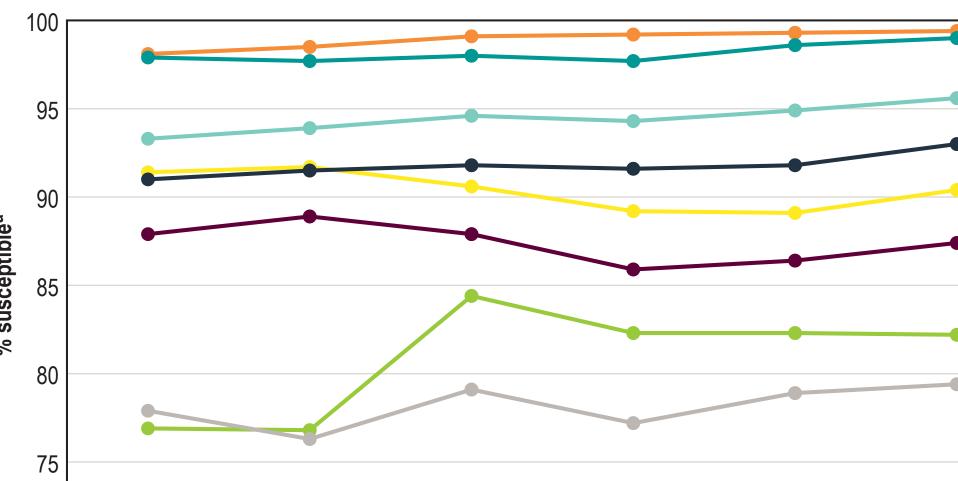
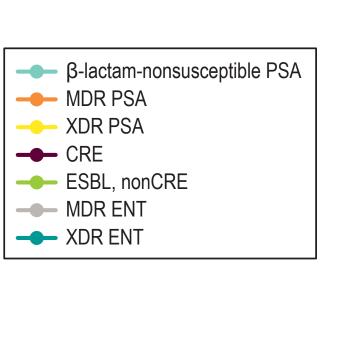


Figure 4 Susceptibility of *Enterobacteriaceae* by year



70							
70	2012 (n=3,633)	2013 (n=4,051)	2014 (n=3,530)	2015 (n=3,948)	2016 (n=3,694)	2017 (n=3,937)	2018 (n=3,914)
Ceftolozane-tazobactam	93.3	93.9	94.6	94.3	94.9	95.6	95.6
Amikacin	98.1	98.5	99.1	99.2	99.3	99.4	99.3
Cefepime	91.4	91.7	90.6	89.2	89.1	90.4	88.8
Ceftazidime	87.9	88.9	87.9	85.9	86.4	87.4	86.6
Colistin ^b	76.9	76.8	84.4	82.3	82.3	82.2	81.7
Levofloxacin	77.9	76.3	79.1	77.2	78.9	79.4	78.1
Meropenem	97.9	97.7	98.0	97.7	98.6	99.0	99.0
Piperacillin-tazobactam	91.0	91.5	91.8	91.6	91.8	93.0	92.3

^a CLSI (2019). ^b EUCAST (2019).



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(n=910)	2018 (n=938)
7.7	95.8
4.8	93.8
1.4	80.4

94.8	93.8
81.4	80.4
82.0	80.7
0.00	99.6
61.9	62.9
76.0	76.4
78.0	76.4

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••	
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- PSA %S trended lower over time for LEV and β -lactams FEP, CAZ, MEM, and PIP-TAZ
- -For ENT, AMK (98.1–99.4%S), MEM (97.7–99.0%S), and C-T (93.3–95.6%S) were the most active (Figure 4)
- No trends were observed for %S for ENT

Conclusions

- The %S for the agents tested varied <10% over the 7-year period
- The frequency of resistant phenotypes in PSA increased, with % of MDR, XDR, and BL-NS highest in 2018
- CRE, MDR, and XDR enterics remained steady, while ESBL, non-CRE increased 6.3%
- Over the study period, C-T %S remained stable at >93.0% for ENT and >95.0% for PSA
- Against all PSA, C-T had similar %S to AMK and COL – Against MDR and XDR PSA, C-T was more active than AMK.
- Against ENT, C-T was the second most active β -lactam after MEM

Acknowledgements

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