Antimicrobial Activity of Cefepime in Combination with Taniborbactam Against Resistant Clinical Isolates from the United States 2018-2021



IHMA, Inc. 2122 Palmer Drive Schaumburg, IL 60173 USA www.ihma.com

M. Hackel¹, M. Wise¹, D. Sahm¹

INTRODUCTION

Taniborbactam is a novel, investigational cyclic boronate-based broad-spectrum βlactamase inhibitor with potent and selective direct inhibitory activity against and metallo-β-lactamases (Ambler Classes A, B, C and D) [1]. Taniborbactam restores the activity of cefepime against many difficult to treat organisms, including cephalosporin- and Enterobacterales and *Pseudomonas aeruginosa*. In this study, we evaluated the activity of cefepime-taniborbactam comparator agents against nonsusceptible (NS)/resistant (R) clinical isolates of Enterobacterales and P. aeruginosa from the United States (US) from a 2018-2021 global surveillance study.

METHODS

MICs of cefepime with taniborbactam fixed at 4 µg/mL and comparators were determined using the CLSI reference method [2] against Enterobacterales (n=4,220) and *P. aeruginosa* (n=1,222) from the United States collected in 2018-2021. Quality control (QC) testing was performed each day of testing as specified by the CLSI [2, 3]. Isolates were collected from community and hospital infections from 35 sites. Avibactam was tested at a fixed concentration of 4 µg/mL in combination with ceftazidime, tazobactam was tested at a fixed concentration of 4 μg/mL in combination with ceftolozane and piperacillin, and vaborbactam was tested at a fixed concentration of 8 μg/mL in combination with meropenem [3]. CLSI 2023 breakpoints were applied for this analysis, with the EUCAST 2023 meropenem-vaborbactam breakpoint applied against *P. aeruginosa* [3, 4]. Resistant phenotypes were based on CLSI 2023 breakpoints [3]. As cefepimetaniborbactam breakpoints have not yet been established, the provisional nonresistant breakpoint of ≤16 μg/mL was considered for comparative purposes. Multidrug resistant (MDR) was defined as resistance to at least one agent from ≥3 drug classes based on CLSI 2023 breakpoints.

¹IHMA, Inc., Schaumburg, IL, USA

RESULTS

Figure 1. Distribution of Enterobacterales isolates by species

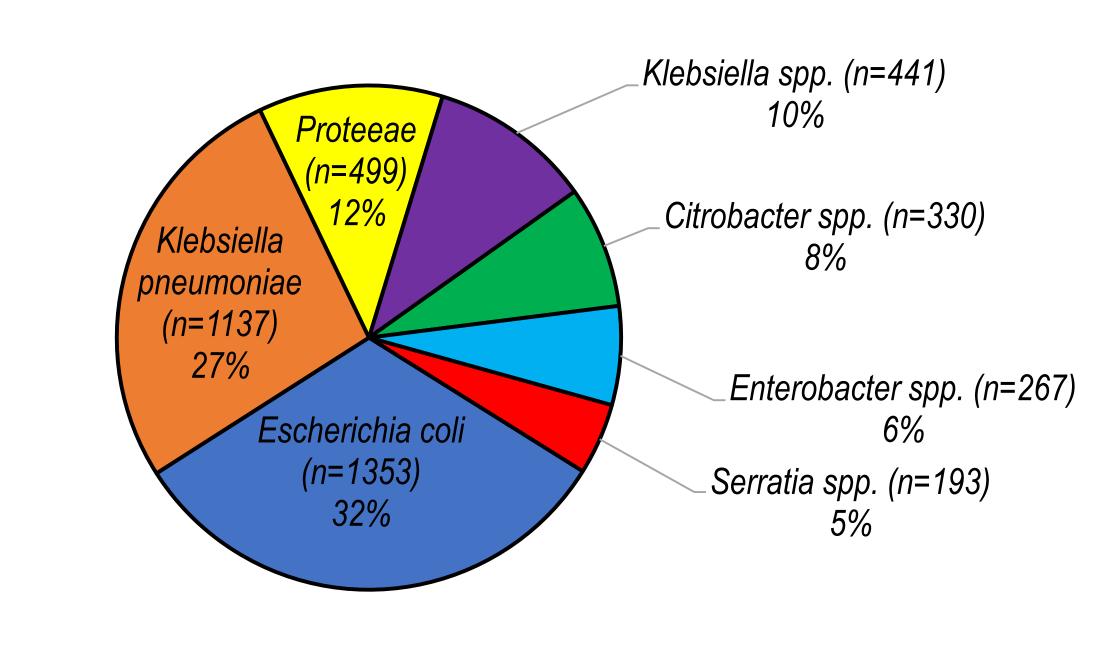
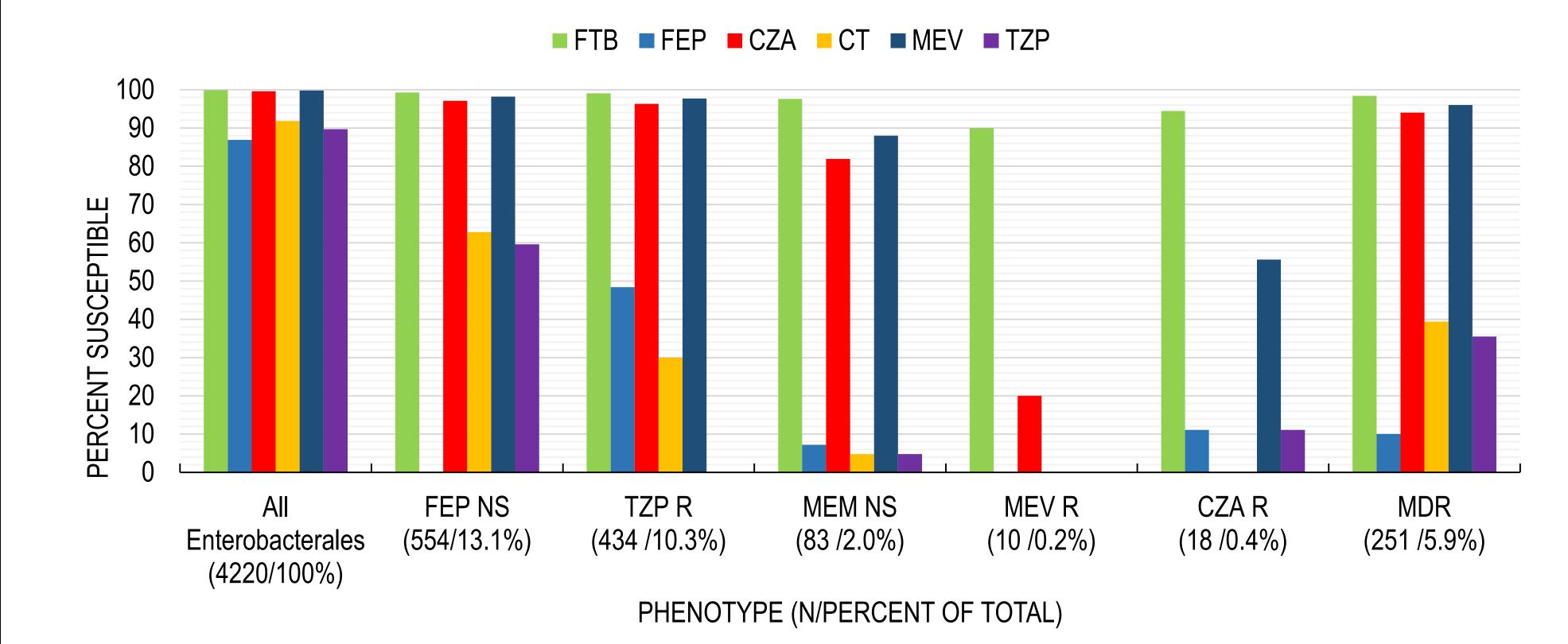


Table 1. Activity of cefepime-taniborbactam and comparators against Enterobacterales

Resistance Phenotype	N (%)	MIC ₉₀ (μg/mL)/Percent susceptible						
		FTB ^a	FEP	CZA	СТ	MEV	TZP	
Enterobacterales	4,220 (100%)	0.12/99.9	8/86.9	0.5/99.6	2/91.8	≤0.06/99.8	32/89.7	
FEP NS	554 (13.1%)	1/99.3	>16/0	2/97.1	>8/62.8	0.12/98.2	>128/59.6	
TZP NS	434 (10.3%)	1/99.1	>16/48.4	2/96.3	>8/30	0.25/97.7	>128/0	
MEM NS	83 (2.0%)	4/97.6	>16/7.2	>16/81.9	>8/4.8	8/88.0	>128/4.8	
MEV NS	10 (0.2%)	8/90.0	>16/0	>16/20	>8/0	>16/0	>128/0	
CZA NS	18 (0.4%)	8/94.4	>16/11.1	>16/0	>8/0	>16/55.6	>128/11.1	
MDR	251 (5.9%)	2/98.4	>16/10.0	4/94.0	>8/39.4	1/96.0	>128/35.5	

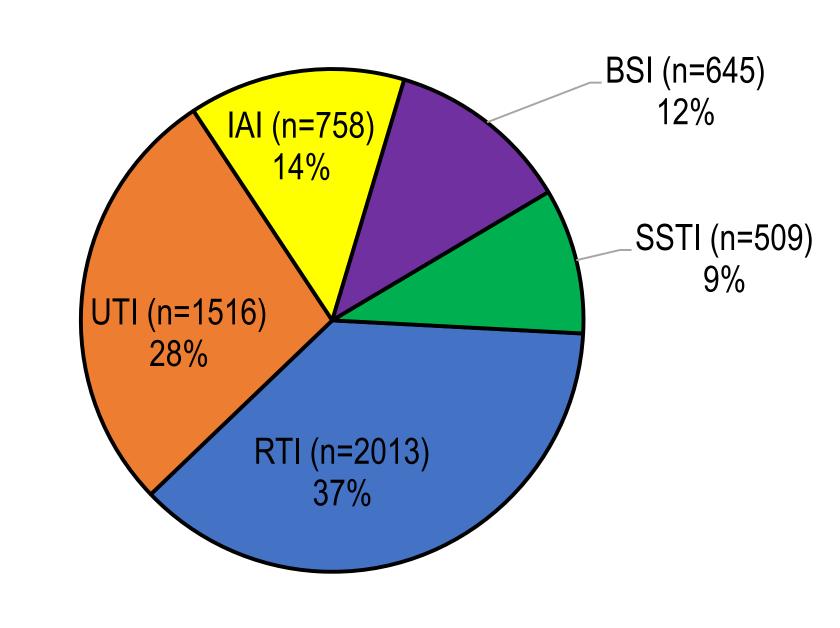
TZP, piperacillin-tazobactam; MDR, multidrug resistant; NS, nonsusceptible; R. resistant. ^aCorresponds to a provisional susceptible breakpoint of ≤16 μg/mL for comparative purposes only

Fig 3. Antimicrobial susceptibility of Enterobacterales overall and by and resistant subset



FTB, cefepime with taniborbactam fixed at 4 µg/mL; FEP, cefepime; CZA, ceftazidime-avibactam; CT, ceftolozane-tazobactam; MEM, meropenem; MEV, meropenem-vaborbactam; TZP, piperacillin-tazobactam; MDR, multidrug resistant; R, resistant; NS, non-susceptible; FTB susceptibility corresponds to a provisional susceptible breakpoint of ≤16 µg/mL for comparative purposes

Figure 2. Distribution of isolates by infection sources



infection: IAI, intraabdominal infection: RTI, respiratory tract infection: SSTI, skin/soft tissue infection UTI, urinary tract infection

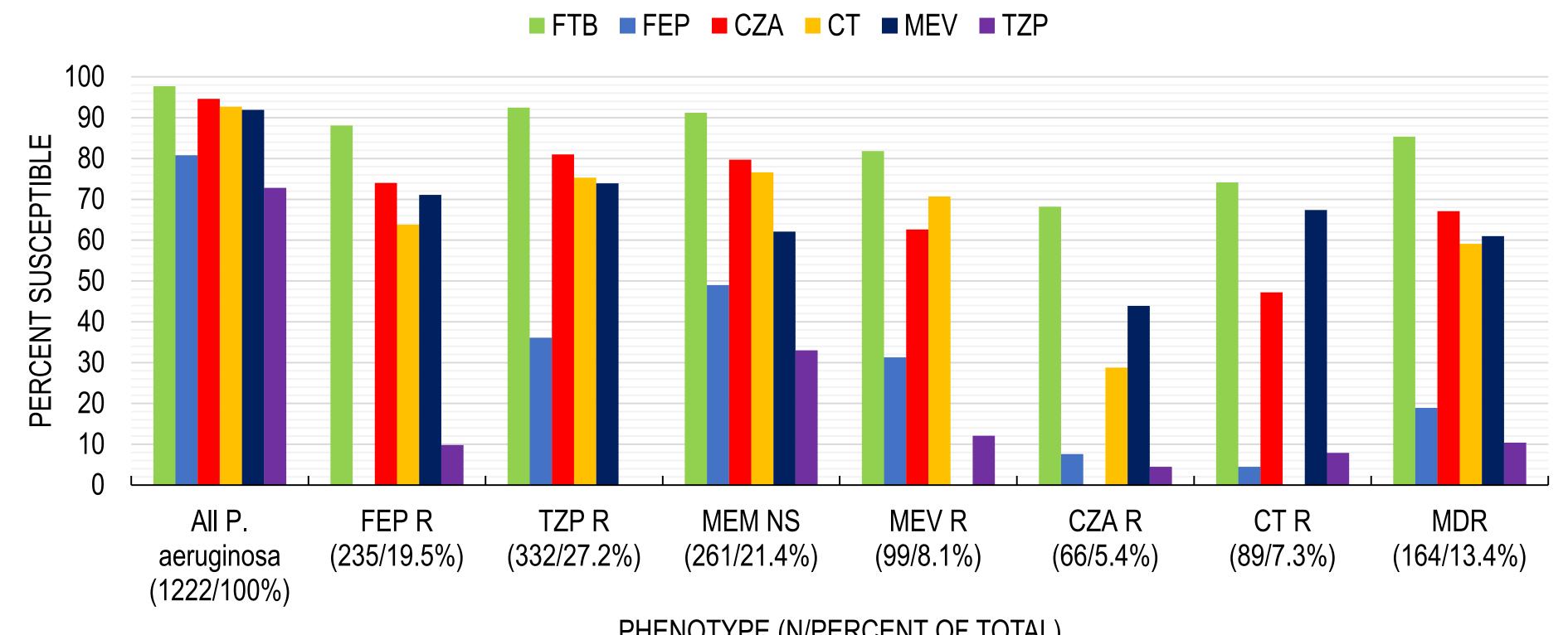
Table 2. Activity of cefepime-taniborbactam and comparators against *P. aeruginosa*

Resistance Phenotype	N (%)		MIC ₉₀ (μg/mL)/Percent susceptible							
		FTB ^a	FEPb	CZA	СТ	MEV	TZPb			
P. aeruginosa	1222/100	8/97.7	32/80.8	8/94.6	4/92.7	8/91.9	>128/72.8			
FEP NS	235/19.2	32/88.1	>32/0	>16/74.0	>16/63.8	>16/71.1	>128/9.8			
TZP NS	332/27.2	16/92.5	>32/36.1	16/81.0	16/75.3	>16/73.9	>128/0			
MEM NS	261/21.4	16/91.2	>32/49.0	>16/79.7	16/76.6	>16/62.1	>128/33.0			
MEV Rb	99/8.1	>32/81.8	>32/31.3	>16/62.6	>16/70.7	>16/0	>128/12.1			
CZA R	66/5.4	>32/68.2	>32/7.6	>16/0	>16/28.8	>16/43.9	>128/4.5			
CT NS	89/7.3	>32/74.2	>32/4.5	>16/47.2	>16/0	>16/67.4	>128/7.9			
MDR	164/13.4	32/85.4	>32/18.9	>16/67.1	>16/59.1	>16/61.0	>128/10.4			

FTB, cefepime with taniborbactam fixed at 4 µg/mL; FEP, cefepime; CZA, ceftazidime-avibactam; CT, ceftolozane-tazobactam; MEM, meropenem; MEV, meropenem-vaborbactam; TZP. piperacillin-tazobactam; MDR, multidrug resistant; NS, nonsusceptible; R, resistant.

^aCorresponds to a provisional susceptible breakpoint of ≤16 µg/mL for comparative purposes on bEUCAST breakpoints (susceptible ≤8 μg/mL/resistant ≥16 μg/mL) applied for *P. aeruginosa*

Fig 4. Antimicrobial susceptibility of *P. aeruginosa* overall and by and resistant subsets



PHENOTYPE (N/PERCENT OF TOTAL)

FTB, cefepime with taniborbactam fixed at 4 µg/mL; FEP, cefepime; CZA, ceftazidime-avibactam; CT, ceftolozane-tazobactam; MEM, meropenem; MEV, meropenem-vaborbactam TZP, piperacillin-tazobactam; MDR, multidrug resistant; R, resistant; MEV R based on EUCAST 2023 breakpoint; NS, non-susceptible; FTB susceptibility corresponds to a provisional susceptible breakpoint of ≤16 μg/mL for comparative purposes

RESULTS SUMMARY

- E. coli (32% of total isolates) and K. pneumoniae (27% of total isolates) were the most common species of Enterobacterales represented in this surveillance collection (Figure 1).
- Overall, 13.1% and 10.3% of Enterobacterales isolates from the United States were nonsusceptible to cefepime and piperacillin-tazobactam, respectively (Table 1). A total of 5.9% of isolates were MDR, and 2.0% were nonsusceptible to meropenem. K. pneumoniae (n=1,137) accounted for 37.2, 31.3, 60.2, and 41.4% of cefepime nonsusceptible, piperacillin-tazobactam nonsusceptible, MDR, and meropenem nonsusceptible isolates among Enterobacterales species within each subset, respectively.
- Cefepime-taniborbactam had potent activity against Enterobacterales overall, with an MIC₉₀ value of 0.12 μg/mL and 99.9% inhibited at ≤16 μg/mL (Table 1, Figure 2).
- Cefepime-taniborbactam maintained activity against resistant subsets of Enterobacterales (MIC₉₀ range, 1 to 8 μ g/mL; 90.0% to 99.3% inhibited at ≤16 μ g/mL) including MDR isolates (MIC₉₀, 2 μ g/mL; 98.4% inhibited at ≤16 µg/mL) (Table 1, Figure 1). Greater than 90% of isolates that were nonsusceptible to ceftazidime-avibactam and/or meropenem-vaborbactam were inhibited at ≤16 µg/mL cefepimetaniborbactam.
- From 19.2% to 27.2% of *P. aeruginosa* isolates were nonsusceptible to cefepime, piperacillin-tazobactam and/or meropenem (Table 2). Between 5.4% and 8.1% of isolates were nonsusceptible/resistant to ceftolozane-tazobactam, ceftazidime-avibactam and/or meropenem-vaborbactam.
- Cefepime-taniborbactam was the most active tested agent against P. aeruginosa overall, with an MIC₉₀ value of 8 μg/mL and 97.7% inhibited at ≤16 μg/mL (Table 2, Figure 3).
- Percentages of P. aeruginosa isolates in the nonsusceptible subsets that were inhibited by ≤16 μg/mL cefepime-taniborbactam ranged from 68.2% for ceftazidime-avibactam resistant isolates to 92.5% for piperacillin-tazobactam nonsusceptible isolates. These compared to 0% to 81.0% susceptible to comparators (Table 2, Figure 3).
- Against MDR P. aeruginosa (13.4% of total isolates), cefepime-taniborbactam maintained activity, with 85.4% of isolates inhibited at ≤16 µg/mL, a substantially greater percentage than the most active comparators, ceftazidime-avibactam (67.1% susceptible), meropenem-vaborbactam (61.0% susceptible), and ceftolozane-tazobactam (59.1% susceptible) (Table 2, Figure 3).

CONCLUSIONS

- Cefepime-taniborbactam demonstrated potent in vitro activity against recent Enterobacterales and P. aeruginosa from the United States, including MDR isolates and isolates nonsusceptible to cefepime, meropenem, piperacillin-tazobactam, ceftazidime-avibactam, ceftolozane-tazobactam, and/or meropenem-vaborbactam.
- These data support continued development of cefepime-taniborbactam as a potential treatment option for challenging infections due to resistant Gram-negative pathogens.

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DISCLOSURES

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