

# Activity of Ceftolozane/Tazobactam against Gram-Negative Isolates from Lower Respiratory Tract Infections – SMART United States 2018

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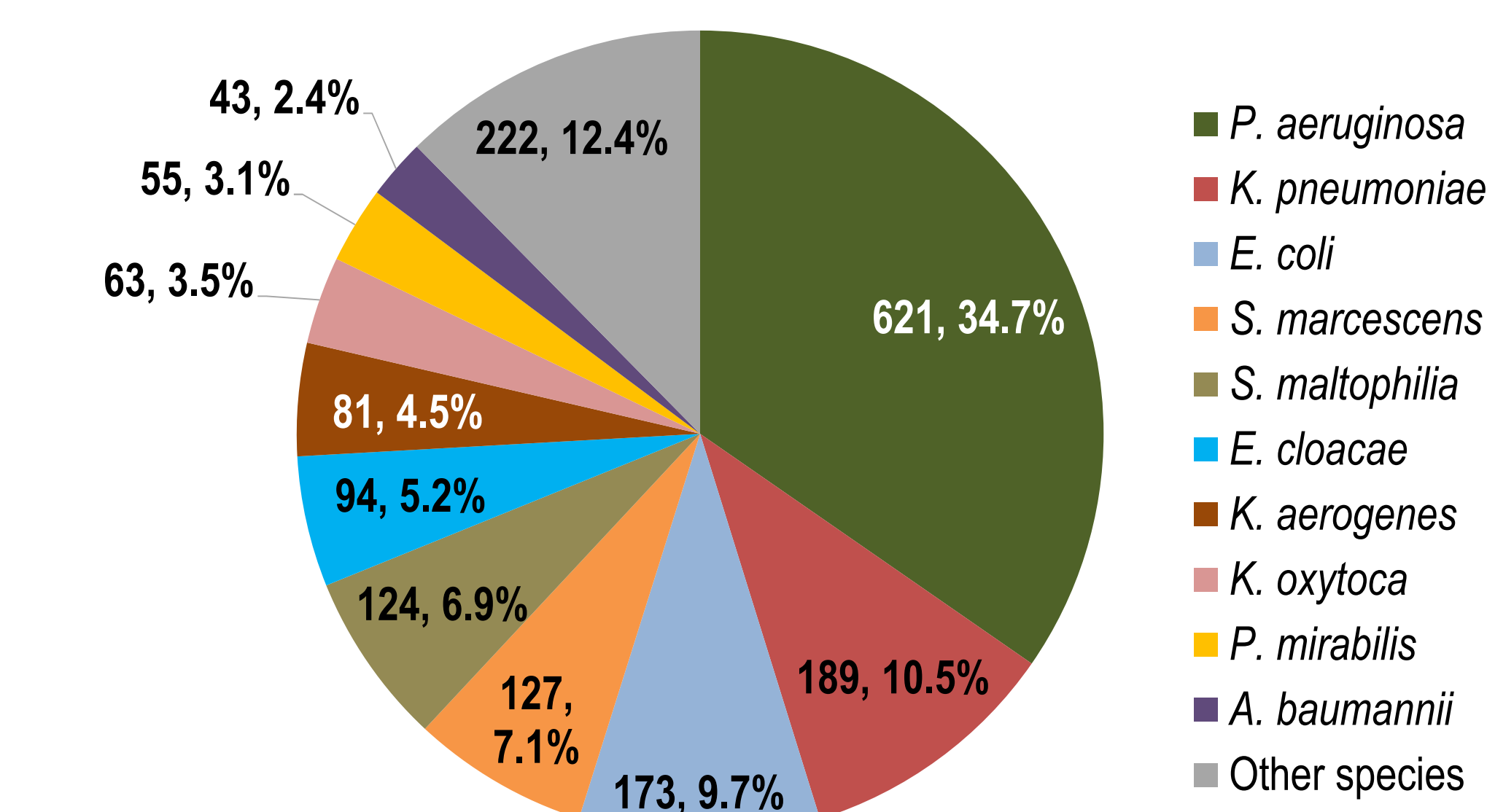
## Introduction

Ceftolozane / tazobactam (C/T) is an anti-pseudomonal cephalosporin combined with a  $\beta$ -lactamase inhibitor. C/T has been approved by the United States Food & Drug Administration (FDA) and the European Medicines Agency (EMA) for complicated urinary tract infections, complicated intraabdominal infections, and hospital-acquired and ventilator-associated bacterial pneumonia. Using isolates collected in the United States as part of the global Study for Monitoring Antimicrobial Resistance Trends (SMART) surveillance program, we evaluated the activity of C/T and comparators against gram-negative pathogens collected from patients with lower respiratory tract infections (LRTI).

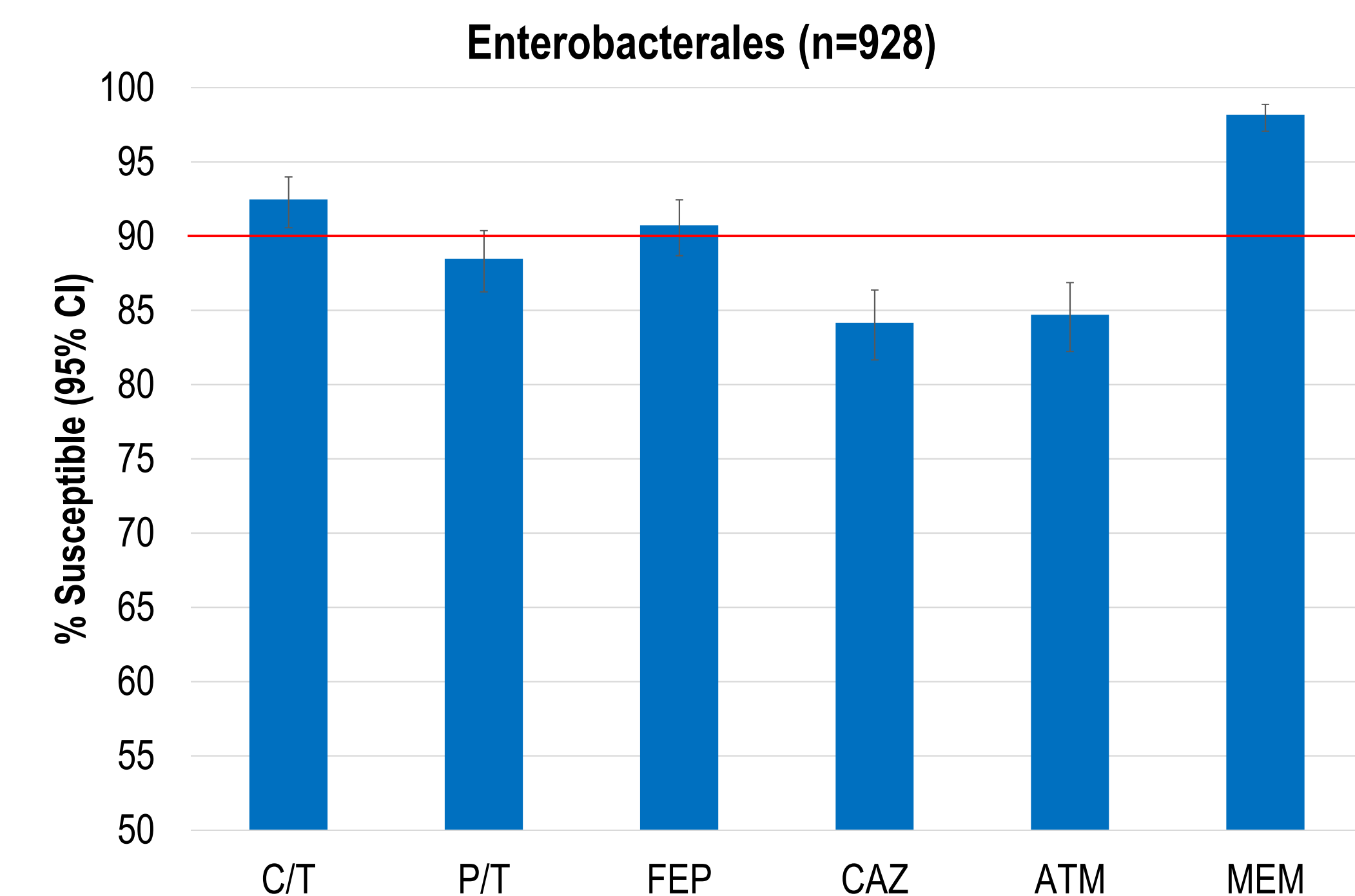
## Methods

In 2018, 24 hospitals in the United States each collected up to 100 consecutive aerobic or facultatively anaerobic gram-negative bacilli from LRTI, for a total of 1792 isolates. MICs were determined using CLSI broth microdilution and interpreted with CLSI breakpoints [1, 2]. C/T-nonsusceptible (NS) Enterobacteriales and *Pseudomonas aeruginosa* isolates were screened by PCR and sequencing for genes encoding  $\beta$ -lactamases [3].

**Figure 1. Species distribution (n, %) among collected gram-negative isolates (n=1792) from patients with LRTI**



**Figure 2. Susceptibility to C/T and  $\beta$ -lactam comparators of all Enterobacteriales combined**



C/T, ceftolozane/tazobactam; P/T, piperacillin/tazobactam; FEP, cefepime; CAZ, ceftazidime; ATM, aztreonam; MEM, meropenem; CI, confidence interval.

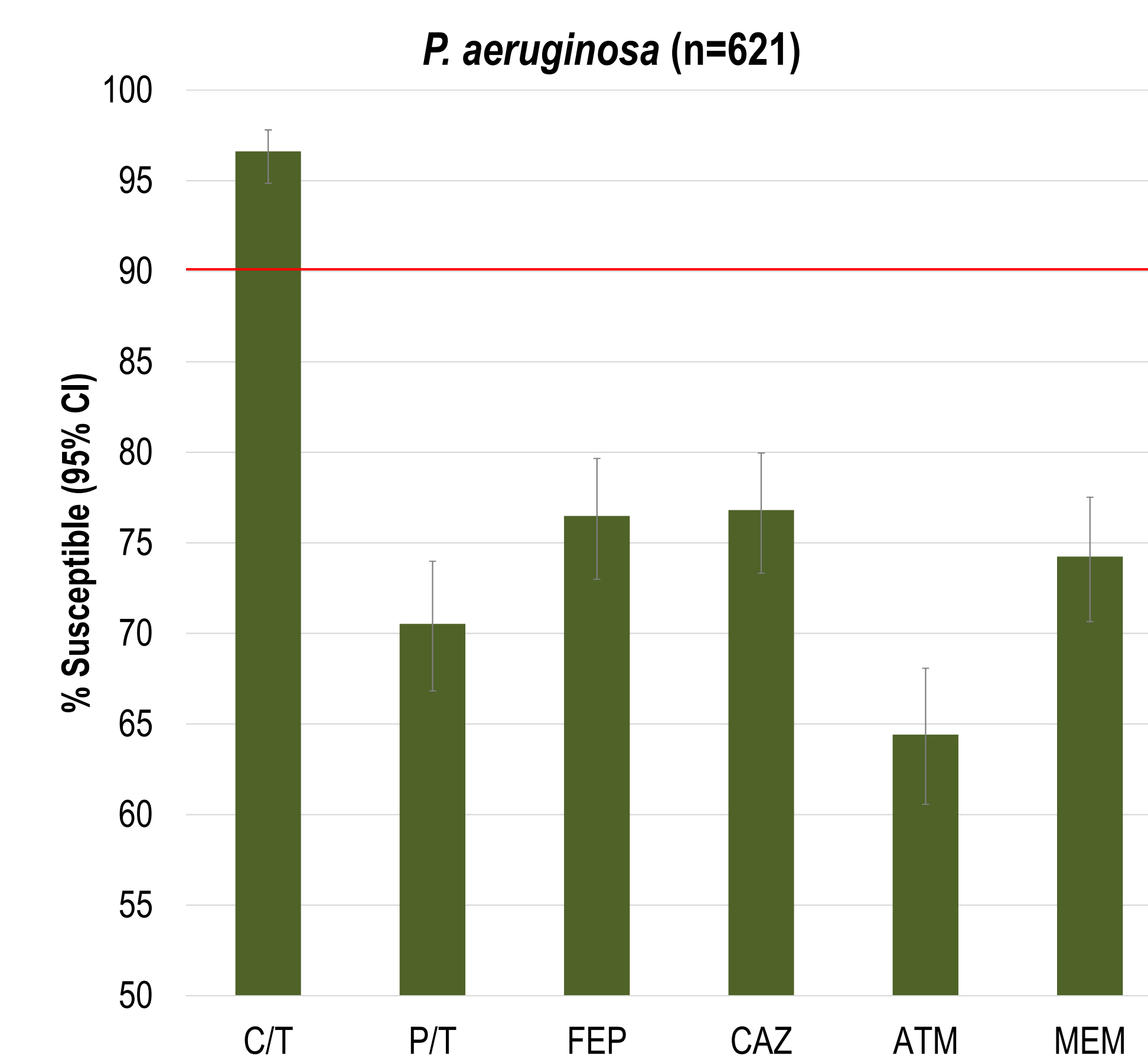
**Table 1. Susceptibility to C/T and  $\beta$ -lactam comparators of the most common Enterobacteriales species**

Species	n	% Susceptible					
		C/T	P/T	FEP	CAZ	ATM	MEM
<i>K. pneumoniae</i>	189	89.9	82.5	78.8	75.7	77.8	95.2
<i>E. coli</i>	173	98.3	92.5	89.0	85.6	85.0	100
<i>S. marcescens</i>	127	96.1	93.7	96.9	95.3	92.9	96.1
<i>E. cloacae</i>	94	85.1	86.2	90.4	76.6	77.7	98.9

C/T, ceftolozane/tazobactam; P/T, piperacillin/tazobactam; FEP, cefepime; CAZ, ceftazidime; ATM, aztreonam; MEM, meropenem

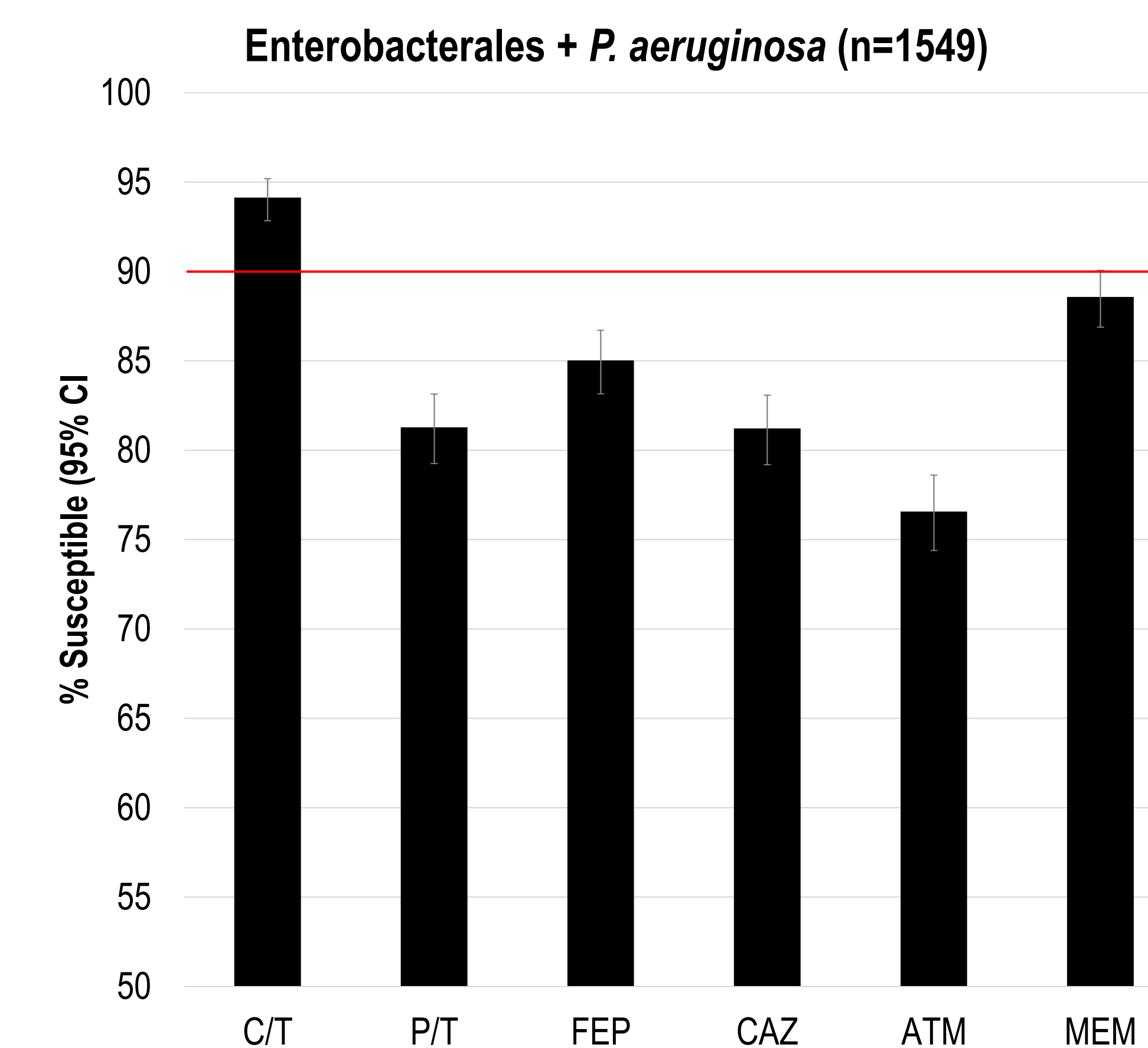
## Results

**Figure 3. Susceptibility to C/T and  $\beta$ -lactam comparators of *P. aeruginosa***



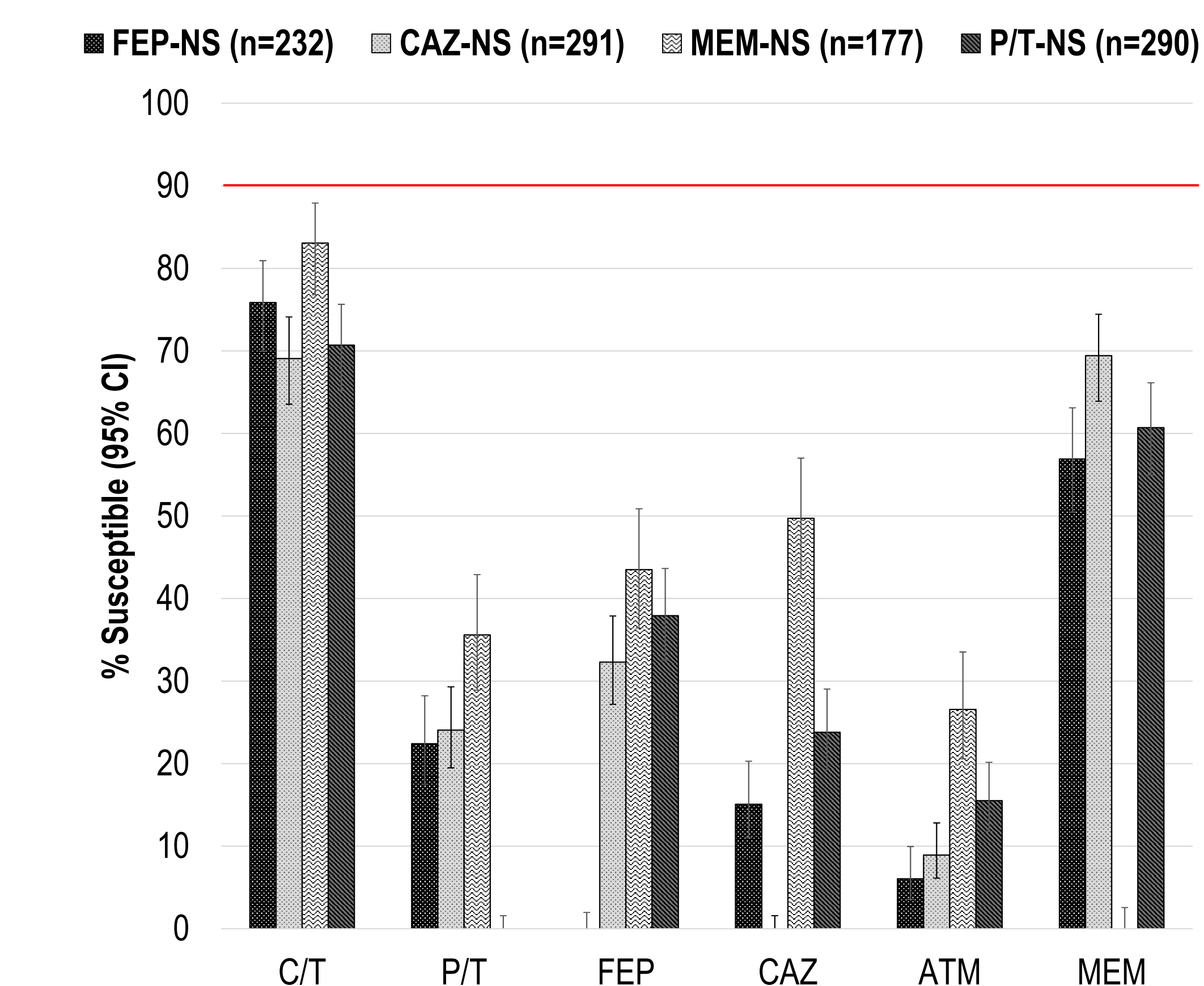
C/T, ceftolozane/tazobactam; P/T, piperacillin/tazobactam; FEP, cefepime; CAZ, ceftazidime; ATM, aztreonam; MEM, meropenem; CI, confidence interval

**Figure 4. Susceptibility to C/T and  $\beta$ -lactam comparators of Enterobacteriales and *P. aeruginosa* combined**



C/T, ceftolozane/tazobactam; P/T, piperacillin/tazobactam; FEP, cefepime; CAZ, ceftazidime; ATM, aztreonam; MEM, meropenem; CI, confidence interval

**Figure 5. Susceptibility to C/T and  $\beta$ -lactam comparators of nonsusceptible phenotypes of Enterobacteriales and *P. aeruginosa* combined**



C/T, ceftolozane/tazobactam; P/T, piperacillin/tazobactam; FEP, cefepime; CAZ, ceftazidime; ATM, aztreonam; MEM, meropenem; CI, confidence interval

**Table 2. Acquired  $\beta$ -lactamases detected in molecularly characterized C/T-nonsusceptible Enterobacteriales and *P. aeruginosa* isolates<sup>a, b</sup>**

Genotype	n (%)	
	Enterobacteriales (n=67)	<i>P. aeruginosa</i> (n=21)
KPC $\pm$ ESBL $\pm$ AmpC	13 (19.4%)	
IMP		1 (4.8%)
AmpC $\pm$ ESBL	1 (1.5%)	
ESBL only	11 (16.4%)	
None detected	42 (62.7%) <sup>c</sup>	20 (95.2%) <sup>d</sup>

<sup>a</sup>Original spectrum  $\beta$ -lactamases (e.g., TEM-1, SHV-1) and intrinsic AmpC  $\beta$ -lactamases common to *P. aeruginosa* and some Enterobacteriales species such as Enterobacter are not included in this analysis.

<sup>b</sup>Three C/T-NS Enterobacteriales isolates were not molecularly characterized.

<sup>c</sup>Among the 42 Enterobacteriales isolates in which no acquired  $\beta$ -lactamases were detected, 39 (92.9%) were species with intrinsic AmpC.

<sup>d</sup>Other resistance mechanisms such as AmpC subtypes with mutations in the  $\Omega$ -loop or in amino acids that interact with it, or undetected  $\beta$ -lactamases may be involved [4]. ESBL, extended-spectrum  $\beta$ -lactamase.

## Results Summary

- Among all gram-negative pathogens collected from patients with LRTI, the 3 most common species were *P. aeruginosa*, *K. pneumoniae*, and *E. coli* (Figure 1). Enterobacteriales and *P. aeruginosa* combined comprised 86% of all collected gram-negative isolates from LRTI.
- C/T was active against 92.5% of all Enterobacteriales isolates combined. Among the comparator  $\beta$ -lactams, only meropenem showed higher activity (Figure 2).
- C/T was active against 85-98% of the 4 most common Enterobacteriales species (Table 1).
- Among *P. aeruginosa* isolates, susceptibility to C/T was 96.6%, 20-32 percentage points higher than to the tested comparator  $\beta$ -lactams (Figure 3).
- C/T was active against 94.1% of all Enterobacteriales and *P. aeruginosa* combined, 6-18 percentage points higher than the other tested comparator agents (Figure 4).
- Among subsets of Enterobacteriales and *P. aeruginosa* isolates that were nonsusceptible to commonly used  $\beta$ -lactams, including meropenem, C/T maintained activity against 69-83% of isolates (Figure 5).
- Among molecularly characterized C/T-NS Enterobacteriales isolates, 19% carried KPC, 16% carried ESBL, and 58% were species with intrinsic AmpC in which no acquired  $\beta$ -lactamases were detected. Among 21 molecularly characterized C/T-NS *P. aeruginosa*, one isolate carried an IMP-type metallo- $\beta$ -lactamase, and in the remaining isolates no acquired  $\beta$ -lactamases were detected (Table 2).

## Conclusions

With its broad coverage of Enterobacteriales and *P. aeruginosa*, C/T can provide an important empiric therapy option for patients with LRTI in the United States, including those with infections caused by meropenem-nonsusceptible isolates.

## References:

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