

Activity of gepotidacin against molecularly characterized *Klebsiella pneumoniae* isolates from patients with urinary tract infections in Europe and adjacent regions (2019–2022)

Gepotidacin demonstrated activity against *K. pneumoniae* carrying β -lactamase genes, including serine carbapenemases, such as *bla*_{KPC} and *bla*_{OXA-48} variants, and metallo- β -lactamase genes.



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Introduction

- Gepotidacin is a novel, bactericidal, first-in-class triazaacenaphthylene antibiotic that inhibits bacterial DNA replication by a unique mechanism of action and distinct binding site, providing a well-balanced inhibition of two different Type II topoisomerase enzymes.
- Gepotidacin has shown *in vitro* activity against most strains of target pathogens, such as *Escherichia coli*, *Staphylococcus saprophyticus*, and *Neisseria gonorrhoeae*, including those resistant to current antibiotics.
- Results from two phase 3 clinical trials demonstrated the efficacy of gepotidacin for the treatment of uncomplicated urinary tract infections (uUTIs). More recently, gepotidacin met its primary efficacy endpoint of non-inferiority in a phase 3 trial comparing gepotidacin with intramuscular ceftriaxone plus oral azithromycin combination for the treatment of urogenital gonorrhoea.
- This study reports on the *in vitro* activity of gepotidacin and other oral antibiotics against molecularly characterized *Klebsiella pneumoniae* collected from patients with UTI in Europe.

Materials and Methods, continued

Screening of resistance determinants

- K. pneumoniae* with MIC results ≥ 2 mg/L for aztreonam and/or ceftazidime and/or ceftriaxone were defined as presumptive ESBL producers and subjected to genome sequencing followed by β -lactamase gene screening.
- Selected isolates had total genomic DNA extracted by the fully automated Thermo Scientific™ KingFisher™ Flex Magnetic Particle Processor (Cleveland, OH, USA), which was used to generate input material for library construction.
- DNA libraries were prepared using the Nextera™ library or Illumina DNA Prep construction protocol (Illumina, San Diego, CA, USA) following the manufacturer's instructions and were sequenced on MiSeq or NextSeq Sequencer platforms at JMI Laboratories.
- FASTQ format sequencing files for each sample set were assembled independently using the *de novo* assembler SPAdes 3.11.0. An in-house software was applied to align the assembled sequences against a comprehensive in-house database containing known β -lactamase genes.

Materials and Methods

Bacterial organisms

- A total of 807 *K. pneumoniae* causing UTI during 2019–2022 in 38 sites in 17 European countries, Israel and Türkiye were included as part of the gepotidacin uropathogen global surveillance study.
- Bacterial identification was confirmed by standard algorithms supported by matrix-assisted laser desorption ionization-time of flight mass spectrometry (Bruker Daltonics, Bremen, Germany).

Susceptibility testing

- Isolates were tested for susceptibility by broth microdilution and agar dilution following Clinical and Laboratory Standards Institute (CLSI) M07 guidelines.
- Interpretation of MIC results was performed using the European Committee on Antimicrobial Susceptibility Testing (EUCAST) criteria, except for amoxicillin-clavulanate MIC values that were interpreted using CLSI breakpoints.

Results

- Gepotidacin had MIC₅₀ and MIC₉₀ values of 4 mg/L and 8 mg/L, respectively, against isolates that did not meet the MIC criteria for screening of β -lactamases (Table 1).
 - Other oral comparator agents showed susceptibility rates of 88.9–95.6% against the group of isolates that did not meet the MIC criteria for screening of β -lactamases.
- A total of 40.3% (323/801) *K. pneumoniae* were selected for screening of β -lactamases.
- The majority of these isolates carried *bla*_{CTX-M} alone (64.7%; 209/323) and a small number carried pAmpC alone (3.1%; 10/323) or other ESBL genes or combinations (6.2%; 20/239).
 - Among β -lactamase screened isolates, 22.6% (73/323) carried carbapenemases, and these isolates originated mostly from Greece (11/73, 15.1%), Italy (9/73, 12.3%), Spain (9/73, 12.3%) and Türkiye (17/73, 23.3%) (data not shown).
- Gepotidacin had an MIC₉₀ of 32 mg/L against strains that meet the MIC criteria for screening of β -lactamases, regardless of β -lactamase gene detected, including carbapenemases.
 - Other oral agents had limited activity (0–60% susceptible) against strains carrying β -lactamase genes, except for mecillinam with susceptibility rates of 88.0–95.0%.

Table 1. Activity of gepotidacin and comparator agents tested against *K. pneumoniae* subsets from Europe and adjacent regions (2019–2022)

Phenotype/genotype (No. isolates)	MIC ₅₀ /MIC ₉₀ in mg/L (% susceptible by EUCAST criteria)					
	GEP	A/C	CFZ	CIP	MEC	SXT
Not-screened (478)	4/8 (—)	2/8 (91.6)	1/4 (91.4)	0.015/0.5 (88.9)	0.25/2 (95.6)	$\leq 0.12/4$ (89.5)
Screened ^a (323)	8/32 (—)	16/>32 (22.3)	>32/>32 (0.3)	>4/>4 (18.0)	4/>32 (75.2)	>4/>4 (19.2)
ESBL/pAmpC (239)	8/32 (—)	16/32 (29.7)	>32/>32 (0.8)	4/>4 (18.0)	2/16 (88.7)	>4/>4 (14.6)
CTX-M ^b (209)	8/32 (—)	16/32 (30.6)	>32/>32 (1.0)	>4/>4 (15.8)	2/16 (88.0)	>4/>4 (11.5)
pAmpC ^c (10)	8/32 (—)	32/>32 (0.0)	>32/>32 (0.0)	0.5/>4 (20.0)	0.5/4 (90.0)	$\leq 0.12/>4$ (60.0)
Other ^d (20)	16/32 (—)	16/>32 (35.0)	>32/>32 (0.0)	0.5/>4 (40.0)	4/8 (95.0)	>4/>4 (25.0)
Carbapenemase ^e (74)	8/32 (—)	>32/>32 (1.4)	>32/>32 (0.0)	>4/>4 (8.2)	>32/>32 (37.0)	>4/>4 (21.9)
Negative ^f (11)	8/32 (—)	16/32 (9.1)	>32/>32 (9.1)	0.03/0.5 (81.8)	16/>32 (36.4)	$\leq 0.12/0.5$ (100)

ESBL, extended-spectrum β -lactamase; GEP, gepotidacin; A/C, amoxicillin-clavulanate; CFZ, cefazolin; CIP, ciprofloxacin; MEC, mecillinam; SXT, trimethoprim-sulfamethoxazole; EUCAST breakpoints and interpretive criteria applied, except for amoxicillin-clavulanate (tested at 2/1 ratio and interpreted per CLSI guidelines); "—" breakpoint not available.

^a Includes isolates with aztreonam, ceftazidime, ceftriaxone or meropenem MICs of ≥ 2 mg/L.

^b Includes 4 *bla*_{CTX-M15}, 3 *bla*_{CTX-M14}, 5 *bla*_{CTX-M16}, 192 *bla*_{CTX-M15}, 1 *bla*_{CTX-M55}, 1 *bla*_{CTX-M27}, 1 *bla*_{CTX-M3} and *bla*_{CTX-M15}, 2 *bla*_{CTX-M9} and *bla*_{CTX-M15}.

^c Includes plasmid pAmpC-9 *bla*_{AmpC} and 1 *bla*_{AmpC}.

^d Includes 4 *bla*_{SHV2}, 1 *bla*_{CTX-M15} and *bla*_{SHV2}, 3 *bla*_{CTX-M15} and *bla*_{SHV1}, 1 *bla*_{SHV2} and *bla*_{SHV1}, 1 *bla*_{SHV2} and *bla*_{SHV1} (ESBL SHV variant).

^e Includes 28 *bla*_{OXA-48}, 25 *bla*_{KPC}, 15 metallo- β -lactamases and 6 isolates carrying 2 carbapenemase genes (2 *bla*_{KPC} plus *bla*_{OXA-48} and 4 *bla*_{OXA-48} plus *bla*_{OXA-48}).

^f Includes isolates where pAmpC, ESBL or carbapenemase genes were not detected.

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Conclusions

- Gepotidacin demonstrated activity against *K. pneumoniae* causing UTI in European countries and adjacent regions, including isolates carrying ESBL, pAmpC and/or carbapenemase genes.
- These data support further clinical development of gepotidacin as a potential treatment option for uUTI caused by *K. pneumoniae* including when other oral treatment options showed limited activity due to drug resistance.

Disclosures

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