

In vitro activity of ceftibiprole against clinical isolates collected from blood and respiratory specimen of hospitalized patients: results of the PEG study

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Introduction and purpose

- The selection of initial empirical antibiotic therapy for patients with community-acquired pneumonia (CAP) or hospital-acquired pneumonia (HAP) is based on local antimicrobial profiles and sensitivity patterns and on risk factors for multidrug-resistant pathogens (e.g. previous antibiotic use and recent hospitalization).^{1,2}
- The empirical treatment of HAP has become increasingly complicated by the emergence and spread of methicillin-resistant *Staphylococcus aureus* (MRSA) and multidrug-resistant Gram-negative pathogens, while the treatment of CAP is compromised by antibiotic-resistant pneumococci.³
- Ceftibiprole medocartil, a new-generation cephalosporin, was recently approved in Europe for the treatment of CAP and HAP (excluding ventilator-associated pneumonia [VAP]) in adults. Ceftibiprole, the active moiety of ceftibiprole medocartil, has broad-spectrum *in vitro* activity against Gram-positive and Gram-negative pathogens, including MRSA, *Enterobacteriaceae* and *Pseudomonas aeruginosa*.⁴
- In a double-blind, randomized study involving 781 patients, ceftibiprole medocartil (500 mg i.v. every 8 hours) was non-inferior to ceftazidime (2 g i.v. every 8 hours) plus linezolid (600 mg i.v. every 12 hours) for clinical cure rates at the test-of-cure visit (77.8% vs 76.2%, excluding VAP, for the clinically evaluable set).⁵
- The objective of this study was to evaluate the *in vitro* activity of ceftibiprole against key pathogens associated with HAP and CAP, namely *S. aureus*, *Streptococcus pneumoniae*, *Enterobacteriaceae* and *P. aeruginosa*, collected during a resistance surveillance study conducted by the Paul Ehrlich Society (PEG) in 2010.

Methods

- Isolates from the respiratory tract and blood of hospitalized patients were prospectively collected from 25 laboratories in Germany (n = 21), Switzerland (n = 3) and Austria (n = 1).
 - For each isolate, the patient's age and sex, details of the hospital department and ward (intensive care unit or general ward), specimen type and collection date were recorded.
 - Isolates were shipped to a central laboratory (Antii Infectives Intelligence, Rheinbach, Germany) for re-identification and susceptibility testing.
- Isolates were identified using standard laboratory methods and matrix-assisted laser desorption/ionization time-of-flight mass spectrometry.
- Minimum inhibitory concentrations (MICs) of ceftibiprole and comparators were determined for each isolate by the broth microdilution method according to the international standard ISO 20776-1.⁶ The resulting MICs were interpreted (when possible) using the species-related clinical breakpoints defined by the European Committee on Antimicrobial Susceptibility Testing (Table 1).^{7,8}
- The Clinical and Laboratory Standards Institute MIC method was used to screen for and confirm isolates with an extended-spectrum β -lactamase (ESBL)-producing phenotype.
- Statistically significant differences in resistance rates were determined by comparing 95% confidence intervals (CIs). Intervals were constructed using the Newcombe-Wilson method without continuity correction.

Results

- Of the 1246 isolates tested, 813 (65.2%) were recovered from respiratory tract specimens and 433 (34.8%) from blood. There were 544 (43.7%) isolates from patients in intensive care units and 702 (56.3%) from patients in general wards.
 - In total, 588 (47.2%) isolates were recovered from patients with HAP and 283 (22.7%) from patients with CAP (unknown source for the remaining 375 [30.1%] patients).

Table 1. Minimum inhibitory concentration breakpoints for ceftibiprole.^{7,8}

Organism	EUCAST breakpoint (mg/L) ^a	
	Susceptible \leq	Resistant >
PK/PD breakpoint	4	4
<i>Enterobacteriaceae</i>	0.25	0.25
<i>Pseudomonas aeruginosa</i>	— ^b	— ^b
<i>Staphylococcus aureus</i>	2	2
<i>Streptococcus pneumoniae</i>	0.5	0.5
All other organisms	— ^c	— ^c

^aSpecies-related breakpoint not defined. EUCAST, The European Committee on Antimicrobial Susceptibility Testing; PD, pharmacodynamics; PK, pharmacokinetics.

Organism/phenotype (n)	Number (cumulative %) of isolates inhibited by ceftibiprole at a concentration (mg/L) of:							S (%)			
	≤ 0.25	0.5	1	2	4	8	16				
<i>Staphylococcus aureus</i> (188)	41 (21.8)	115 (83.0)	3 (84.6)	26 (98.4)	3 ^a (100)			98.4			
	Methicillin-susceptible (158)	41 (25.9)	115 (98.7)	1 (99.4)	1 (100)			100.0			
	Methicillin-resistant (30)		2 (6.7)	25 (90.0)	3 ^a (100)			90.0			
<i>Streptococcus pneumoniae</i> (254)	241 (94.9)	10 (98.8)	3 (100)					98.8			
	Penicillin-susceptible (207)	207 (100)						100.0			
	Penicillin-nonsusceptible (47)	34 (72.3)	10 (93.6)	3 (100)				93.6			
<i>Escherichia coli</i> (179)	143 (79.9)	3 (81.6)	2 (82.7)			2 (83.8)	2 (84.9)	79.9			
	ESBL-negative (146)	141 (96.6)	3 (98.6)	2 (100)				96.6			
	ESBL-positive (33)	2 (6.1)				2 (12.1)	2 (18.2)	6.1			
<i>Klebsiella pneumoniae</i> (108)	86 (79.6)	3 (82.4)		2 (84.3)			2 (18.2)	79.6			
	ESBL-negative (90)	86 (95.6)	3 (98.9)		1 (100)			95.6			
	ESBL-positive (18)			1 (5.6)		1 (11.1)	16 (100)	0			
<i>Klebsiella oxytoca</i> (44)	19 (43.2)	11 (68.2)	4 (77.3)					43.2			
	<i>Enterobacter spp.</i> (89) ^b	66 (74.2)	3 (77.5)	1 (78.7)	3 (82.0)	7 (89.9)	3 (93.3)	1 (94.4)	74.2		
	<i>Serratia spp.</i> (74) ^c	63 (85.1)	5 (91.9)	3 (95.9)	1 (97.3)		1 (98.6)		85.1		
<i>Citrobacter spp.</i> (26) ^d	22 (84.6)	2 (92.3)						84.6			
	<i>Proteaceae</i> (43) ^e	33 (76.7)		1 (79.1)	1 (81.4)	1 (83.7)	2 (88.4)	5 (100)	76.7		
	<i>Pseudomonas aeruginosa</i> (241)		2 (0.8)	10 (5.0)	58 (29.0)	69 (57.7)	48 (77.6)	23 (87.1)	7 (90.0)	NE	
Ceftazidime-susceptible (191)			2 (1.0)	9 (5.8)	58 (36.1)	61 (68.1)	39 (88.5)	15 (96.3)	3 (97.9)	4 (100)	79.6
Ceftazidime-resistant (50)				1 (2.0)		8 (18.0)	9 (36.0)	8 (52.0)	4 (60.0)	20 (100)	NE

^aEtest[®] MICs were 3 mg/L for one isolate and 1–1.5 mg/L for two isolates. ^bIncludes *Enterobacter cloacae* (n = 65), *E. aerogenes* (n = 19) and *E. asburiae* (n = 5). ^cIncludes *Serratia marcescens* (n = 71), *S. liquefaciens* (n = 2) and *S. ureilytica* (n = 1). ^dIncludes *Citrobacter freundii* (n = 13), *C. koseri* (n = 11), *C. braakii* (n = 1) and *C. farmeri* (n = 1). ^eIncludes *Proteus mirabilis* (n = 29), *Morganella morganii* (n = 9) and *P. vulgaris* (n = 5). ESBL, extended-spectrum β -lactamase; EUCAST, The European Committee on Antimicrobial Susceptibility Testing; MIC, minimum inhibitory concentration; NE, not evaluable because EUCAST has not defined a species-related breakpoint; S, susceptible.

- Almost two-thirds of isolates (n = 793 [63.6%]) were recovered from male patients, and the median age of patients was 66 years (range, < 1–100 years).

Table 3. *In vitro* activity of ceftibiprole and comparators against Gram-positive isolates collected from the respiratory tract or blood of hospitalized patients.

Organism/phenotype (n)	Antibiotic concentration (mg/L)			Proportion of isolates that are S/I/R (%) ^a		
	MIC ₅₀	MIC ₉₀	MIC range			
<i>Staphylococcus aureus</i> (188)	Ceftibiprole	0.5	≤ 0.25 –4	98.4/–/1.6		
	Daptomycin	0.5	0.5	100/–/0		
	Levofloxacin	≤ 0.25	> 8	≤ 0.25 –> 8	77.7/0/22.3	
	Erythromycin	0.5	> 32	≤ 0.25 –> 32	75.5/0.5/23.9	
	Linezolid	1	1	≤ 0.25 –2	100/–/0	
	Vancomycin	0.5	1	≤ 0.25 –2	100/–/0	
	Methicillin-susceptible (158)	Ceftibiprole	0.5	0.5	≤ 0.25 –2	100/–/0
Daptomycin		0.5	0.5	0.125–1	100/–/0	
Levofloxacin		≤ 0.25	0.5	≤ 0.25 –> 8	92.4/0/7.6	
Erythromycin		0.5	32	≤ 0.25 –> 32	86.7/0/13.3	
Linezolid		1	1	≤ 0.25 –2	100/–/0	
Vancomycin		0.5	1	≤ 0.25 –2	100/–/0	
Methicillin-resistant (30)		Ceftibiprole	2	2	1–4	90.0/–/10.0
	Daptomycin	0.5	1	0.25–1	100/–/0	
	Levofloxacin	> 8	> 8	4–> 8	0/0/100	
	Erythromycin	> 32	> 32	0.5–> 32	16.7/3.3/80.0	
	Linezolid	1	1	≤ 0.25 –2	100/–/0	
	Vancomycin	0.5	1	0.5–1	100/–/0	
	<i>Streptococcus pneumoniae</i> (254)	Ceftibiprole	≤ 0.25	≤ 0.25	≤ 0.25 –1	98.8/–/1.2
Ceftriaxone		≤ 0.125	≤ 0.125	≤ 0.125 –1	97.2/2.8/0	
Erythromycin		≤ 0.25	32	≤ 0.25 –> 32	79.9/0/20.1	
Levofloxacin		1	1	≤ 0.25 –> 8	98.0/–/2.0	
Penicillin		≤ 0.063	0.25	≤ 0.063 –2	81.5/18.5/0	
Penicillin-susceptible (207)		Ceftibiprole	≤ 0.25	≤ 0.25	≤ 0.25	100/–/0
		Ceftriaxone	≤ 0.125	≤ 0.125	≤ 0.125 –0.5	100/0/0
	Erythromycin	≤ 0.25	4	≤ 0.25 –> 32	88.9/0/11.1	
	Levofloxacin	1	1	≤ 0.25 –2	100/–/0	
	Penicillin-nonsusceptible (47)	Ceftibiprole	≤ 0.25	0.5	≤ 0.25 –1	93.6/–/6.4
		Ceftriaxone	0.25	1	≤ 0.125 –1	85.1/14.9/0
		Erythromycin	8	> 32	≤ 0.25 –> 32	40.4/0/59.6
Levofloxacin		1	8	≤ 0.25 –> 8	89.4/–/10.6	

^aSensitivity categories defined by species-related breakpoints according to EUCAST. EUCAST, The European Committee on Antimicrobial Susceptibility Testing; MIC, minimum inhibitory concentration; MIC₅₀, minimum concentration required to inhibit 50% of isolates; MIC₉₀, minimum concentration required to inhibit 90% of isolates; S, susceptible; I, intermediate; R, resistant.

In vitro activity of ceftibiprole and comparators against Gram-positive isolates

- Of the *S. aureus* isolates (n = 188), 158 (84%) were methicillin-susceptible (MSSA) and 30 (16%) were MRSA (Table 2).
- Ceftibiprole demonstrated potent activity against *S. aureus* (Table 2). MIC₅₀ and MIC₉₀ (MIC_{50/90}) values were 0.5/0.5 mg/L for MSSA isolates and 2/2 mg/L for MRSA isolates (Table 3). Overall, 98.4% of *S. aureus* isolates were susceptible to ceftibiprole.
 - All MSSA isolates and 27/30 (90.0%) of the MRSA isolates were susceptible to ceftibiprole (Table 3).
 - The MICs of the three ceftibiprole-resistant MRSA isolates were 4 mg/L (Table 2). When these isolates were retested by Etest[®], MICs were 3 mg/L (resistant) for one isolate and 1–1.5 mg/L (susceptible) for two isolates, leaving only one truly resistant isolate.
- Rates of resistance to levofloxacin and erythromycin were 7.6% and 13.3%, respectively, for MSSA isolates and 100% and 80%, respectively, for MRSA isolates. All *S. aureus* isolates were susceptible to daptomycin, linezolid and vancomycin (Table 3).
- Of the *S. pneumoniae* isolates (n = 254), 47 (18.5%) showed reduced susceptibility to penicillin, but none was resistant to penicillin (Table 2).
- Ceftibiprole inhibited all *S. pneumoniae* isolates at 1 mg/L, although 3 (1.2%) of these isolates (from blood, with penicillin MICs of 1–2 mg/L) were categorized as ceftibiprole-resistant (Table 2).
- Resistance to erythromycin and levofloxacin was observed in 20.1% and 2.0% of *S. pneumoniae* isolates, respectively (Table 3).

In vitro activity of ceftibiprole and comparators against Gram-negative isolates

- Of the *Enterobacteriaceae* isolates (n = 563), 59 (10.5%) showed an ESBL-positive phenotype, including 33/179 (18.4%) *Escherichia coli* isolates and 18/108 (16.7%) *Klebsiella pneumoniae* isolates.
- In total, 432 (76.7%) *Enterobacteriaceae* isolates were susceptible to ceftibiprole (Table 2). MIC_{50/90} values for ceftibiprole against *E. coli* and *K. pneumoniae* were similar to those for ceftriaxone (Table 4), but ceftibiprole showed greater activity against *Enterobacteriaceae* species that produce chromosomally encoded AmpC-lactamases (*Serratia marcescens* and *Enterobacter cloacae*; data not shown).
- For *E. coli* and *K. pneumoniae*, susceptibility rates to ceftibiprole were slightly lower than those for ceftriaxone (differences not significant; Table 4).
- Similarly to other broad-spectrum cephalosporins, ceftibiprole showed weak activity against isolates with an ESBL-positive phenotype; however, two *E. coli* isolates with an ESBL-positive phenotype were susceptible to ceftibiprole (Table 2).
- For *P. aeruginosa*, the MIC_{50/90} values for ceftibiprole (4/32 mg/L) were similar to those for ceftazidime and cefepime (2/32 mg/L and 4/32 mg/L, respectively; Table 4).

Table 4. *In vitro* activity of ceftibiprole and comparators against isolates of the most frequent Gram-negative species collected from the respiratory tract or blood of hospitalized patients.

Organism/phenotype (n)	Antibiotic concentration (mg/L)			Proportion of isolates that are S/I/R (%) ^a		
	MIC ₅₀	MIC ₉₀	MIC range			
<i>Escherichia coli</i> (179)	Ceftibiprole	≤ 0.25	> 32	≤ 0.25 –> 32	79.9/–/20.1	
	Ceftriaxone	≤ 0.125	> 16	≤ 0.125 –> 16	81.0/1.7/17.3	
	Meropenem	≤ 0.5	≤ 0.5	≤ 0.5	100/0/0	
	Piperacillin-tazobactam	2	64	≤ 1 –> 64	85.5/3.4/11.2	
	Ciprofloxacin	0.125	> 8	≤ 0.063 –> 8	57.0/2.2/40.8	
	Gentamicin	1	> 16	≤ 0.25 –> 16	87.7/0/12.3	
	ESBL-negative (146)	Ceftibiprole	≤ 0.25	≤ 0.25	≤ 0.25 –1	96.6/–/3.4
		Ceftriaxone	≤ 0.125	≤ 0.125	≤ 0.125 –2	97.9/2.1/0
		Meropenem	≤ 0.5	≤ 0.5	≤ 0.5	100/0/0
		Piperacillin-tazobactam	2	16	≤ 1 –> 64	89.7/2.7/7.5
		Ciprofloxacin	≤ 0.063	> 8	≤ 0.063 –> 8	65.8/2.7/31.5
Gentamicin		1	2	≤ 0.25 –> 16	93.2/0/6.8	
ESBL-positive (33)		Ceftibiprole	> 32	> 32	≤ 0.25 –> 32	6.1/–/93.9
		Ceftriaxone	> 16	> 16	0.5–> 16	6.1/0/93.9
		Meropenem	≤ 0.5	≤ 0.5	≤ 0.5	100/0/0
		Piperacillin-tazobactam	2	64	≤ 1 –> 64	66.7/6.1/27.3
		Ciprofloxacin	> 8	> 8	≤ 0.063 –> 8	18.2/0/81.8
	Gentamicin	1	> 16	0.5–> 16	63.6/0/36.4	
	<i>Klebsiella pneumoniae</i> (108)	Ceftibiprole	≤ 0.25	> 32	≤ 0.25 –> 32	79.6/–/20.4
		Ceftriaxone	≤ 0.125	> 16	≤ 0.125 –> 16	84.3/0/15.7
		Meropenem	≤ 0.5	≤ 0.5	≤ 0.5 –1	100/0/0
		Piperacillin-tazobactam	4	32	≤ 1 –> 64	81.5/6.5/12.0
		Ciprofloxacin	≤ 0.063	> 8	≤ 0.063 –> 8	78.7/1.9/19.4
Gentamicin		0.5	8	≤ 0.25 –> 16	88.9/0/11.1	
ESBL-negative (90)		Ceftibiprole	≤ 0.25	≤ 0.25	≤ 0.25 –2	95.6/–/4.4
		Ceftriaxone	≤ 0.125	0.25	≤ 0.125 –0.5	100/0/0
		Meropenem	≤ 0.5	≤ 0.5	≤ 0.5	100/0/0
		Piperacillin-tazobactam	2	8	≤ 1 –> 64	91.1/3.3/5.6
		Ciprofloxacin	≤ 0.063	0.5	≤ 0.063 –> 8	91.1/2.2/6.7
	Gentamicin	0.5	0.5	≤ 0.25 –16	96.7/0/3.3	
	ESBL-positive (18)	Ceftibiprole	> 32	> 32	2–> 32	0/–/100
		Ceftriaxone	> 16	> 16	0.5–> 16	5.6/0/94.4
		Meropenem	≤ 0.5	1	≤ 0.5 –1	100/0/0
		Piperacillin-tazobactam	16	> 64	4–> 64	33.3/22.2/44.4
		Ciprofloxacin	> 8	> 8	≤ 0.063 –> 8	16.7/0/83.3
Gentamicin		1	> 16	≤ 0.25 –> 16	50.0/0/50.0	
<i>Pseudomonas aeruginosa</i> (241)		Ceftibiprole	4	32	0.5–> 32	– ^b /–/–
		Ceftazidime	2	32	0.5–> 32	79.3/–/20.7
		Cefepime	4	32	0.5–> 32	74.7/–/25.3
		Meropenem	1	16	≤ 0.5 –> 32	71.0/14.9/14.1
		Piperacillin-tazobactam	8	64	≤ 1 –> 64	77.2/–/22.8
	Ciprofloxacin	0.25	> 8	≤ 0.063 –> 8	68.0/4.1/27.8	
	Gentamicin	2	16	≤ 0.25 –> 16	86.7/–/13.3	
	Ceftazidime-susceptible (191)	Ceftibiprole	4	16	0.5–> 32	– ^b /–/–
		Ceftazidime	2	4	0.5–8	100/–/0
		Cefepime	4	8	0.5–32	92.1/–/7.9
		Meropenem	≤ 0.5	4	≤ 0.5 –32	84.3/9.5/5.8
Piperacillin-tazobactam		8	16	≤ 1 –64	90.1/–/9.9	
Ciprofloxacin		0.25	4	≤ 0.063 –> 8	78.5/4.2/17.3	
Gentamicin		1	4	≤ 0.25 –> 16	94.8/–/5.2	
Ceftazidime-resistant (50)		Ceftibiprole	16	> 32	1–> 32	– ^b /–/–
		Ceftazidime	32	> 32	16–> 32	0/–/100
		Cefepime	32	> 32	2–> 32	8.0/–/92.0
		Meropenem	8	> 32	≤ 0.5 –> 32	20.0/34.0/46.0
	Piperacillin-tazobactam	64	> 64	8–> 64	28.0/–/72.0	
	Ciprofloxacin	8	> 8	0.125–> 8	28.0/4.0/68.0	
	Gentamicin	4	> 16	≤ 0.25 –> 16	56.0/–/44.0	

^aSensitivity categories defined by species-related breakpoints according to EUCAST. ^bSpecies-related breakpoint not defined. ESBL