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INTRODUCTION

- Ceftolozane-tazobactam is an antibacterial combination of an antipseudomonal cephalosporin and a well-described β-lactamase inhibitor
- Ceftolozane-tazobactam has activity against most common β-lactam resistance mechanisms employed by *Pseudomonas aeruginosa,* including AmpC production (PDC), up-regulated efflux pumps, and porin reductions (OprD loss)
- Ceftolozane-tazobactam also has activity against most ESBL-producing *Enterobacteriaceae* (new taxonomy: Enterobacterales)
- Ceftolozane-tazobactam has been approved in >50 countries, including the US
 Food and Drug Administration in 2014 and the European Medicines Agency
 in 2015 to treat complicated urinary tract infections, acute pyelonephritis, and
 complicated intra-abdominal infections in adults
- Paediatric treatment trials are in progress
- The Program to Assess Ceftolozane-Tazobactam Susceptibility (PACTS) monitors ceftolozane-tazobactam resistance to gram-negative isolates worldwide
- This study analysed the susceptibility of gram-negative isolates from European paediatric patients

MATERIALS AND METHODS

- A total of 2,136 gram-negative isolates were collected during 2012–2017 from paediatric patients (<18 years old) in 50 European hospitals participating in PACTS
- Patients ≤1 year old (yo) contributed 43.9% of isolates
- Other age groups and %: 2-6 yo, 22.0%; 7-12 yo, 18.6%; ≥13 yo, 15.5%
- Isolates were tested for ceftolozane-tazobactam susceptibility (S) by CLSI broth microdilution method at JMI Laboratories using EUCAST breakpoints (2018)
- Other antibiotics tested were amikacin, cefepime, ceftazidime, colistin, levofloxacin, meropenem, and piperacillin-tazobactam
- Ceftolozane-tazobactam and piperacillin-tazobactam were both tested at a fixed 4 mg/L tazobactam concentration
- Antibiotic-resistant phenotypes identified for Enterobacteriaceae (new taxonomy Enterobacterales) included carbapenem-resistant (CRE) and non-CRE extended-spectrum beta-lactamase screen-positive (ESBL, non-CRE)
- Antibiotic-resistant phenotypes identified for *Pseudomonas aeruginosa* included ceftazidime-nonsusceptible, meropenem-nonsusceptible, cefepime-nonsusceptible, piperacillin-tazobactam-nonsusceptible, and nonsusceptible to all 4 β-lactam comparators (β-lactam-nonsusceptible)
- Isolates were considered multidrug-resistant (MDR) if they were nonsusceptible to 3 or more drug classes

RESULTS

- The most common infection type in hospitalised paediatric patients was pneumonia (n=537) followed by bloodstream infection (n=483) and urinary tract infection (n=450), as shown in Figure 1
- The 5 most common species are shown in Figure 2
- The 3 most common species in each infection type were Escherichia coli (n=649), P. aeruginosa (n=425), and Klebsiella pneumoniae (n=287)
- The distribution of these 3 species by infection type is shown in Figure 3

Figure 1 Number of isolates per paediatric infection type in European hospitals (PACTS, 2012–2017)

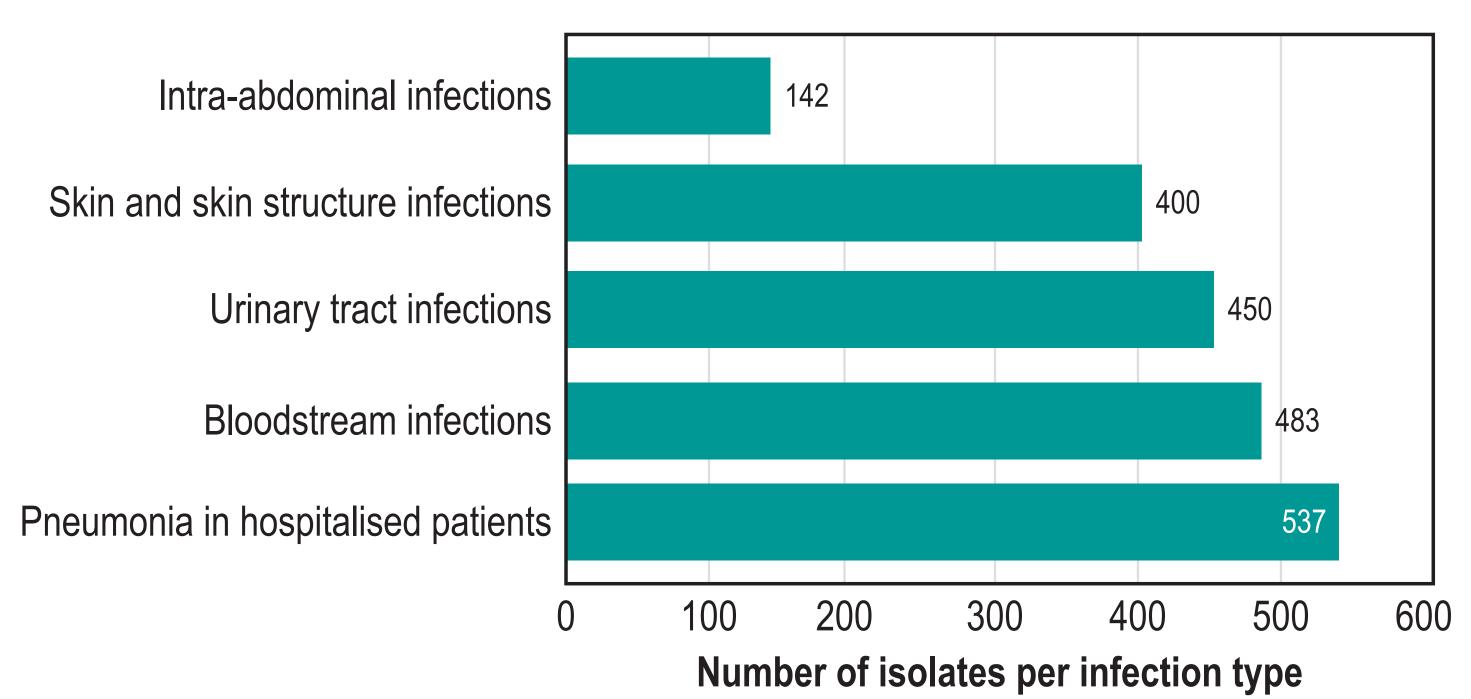
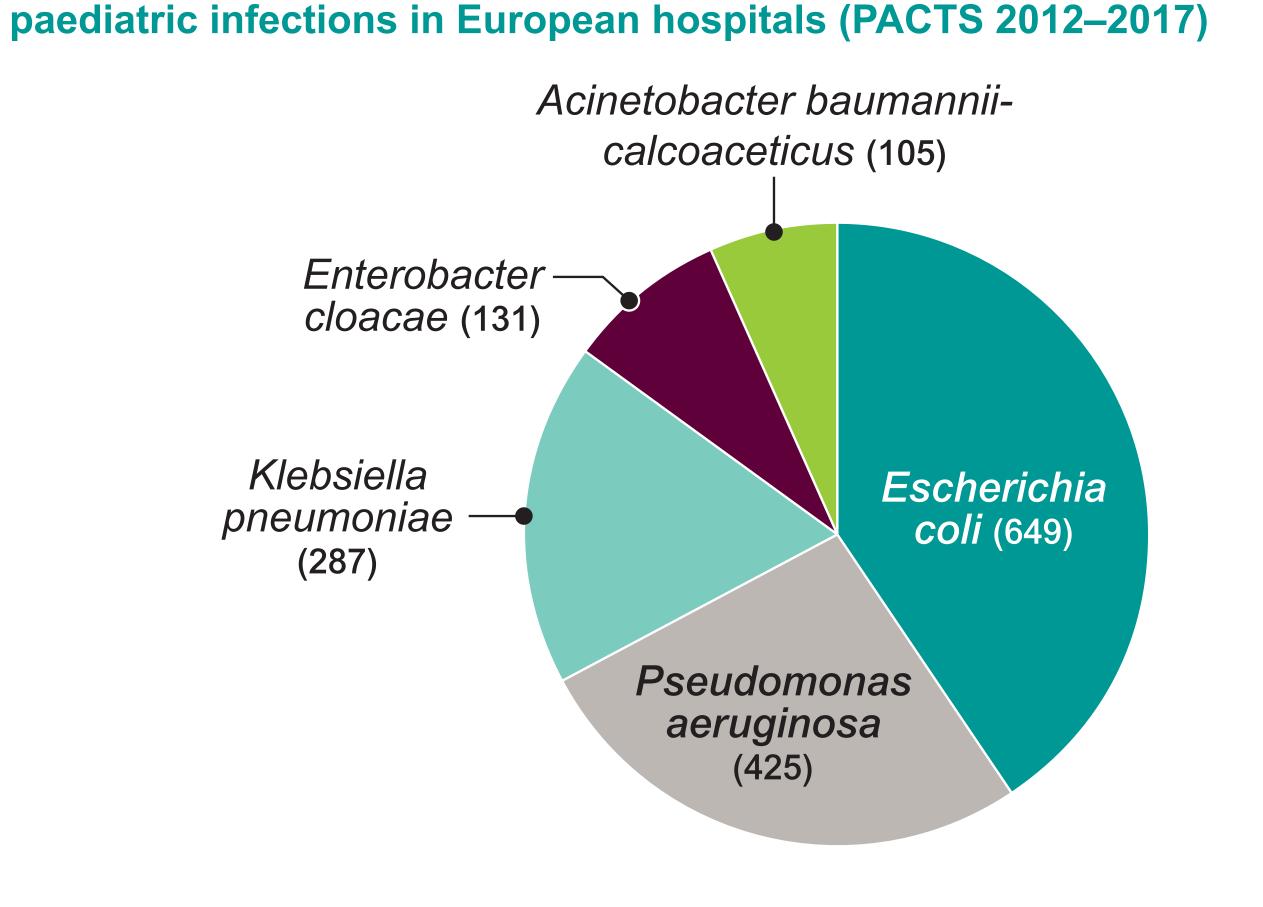


Figure 2 Top 5 pathogens (species and number) isolated from



- E. coli was the most frequent isolate from urinary tract and intraabdominal infections
- P. aeruginosa was the most frequent isolate from pneumonia
- Susceptibilities of ceftolozane-tazobactam and comparators for the main species and resistant phenotypes are shown in Table 1
- For P. aeruginosa, ceftolozane-tazobactam (93.2%S) and colistin (98.4%S)
 were the most active antimicrobials
- The most active drugs against Enterobacteriaceae were meropenem (98.1%S) and amikacin (96.1%S), followed by ceftolozane-tazobactam (89.8%S)
- The ceftolozane-tazobactam MIC distributions for the main pathogens in this study are shown in Table 2

Table 1 Ceftolozane-tazobactam and comparator susceptibility for main pathogen groups and resistant phenotypes (PACTS, 2012–2017)

	n	% susceptible ^a									
Organism/organism group		Ceftolozane- tazobactam	Cefepime	Ceftazidime	Meropenem	Piperacillin- tazobactam	Levofloxacin	Amikacin	Colistin		
Enterobacteriaceae	1,417	89.8	77.0	74.8	98.1	83.0	83.1	96.1	85.8		
ESBL, non-CRE	271	77.1	10.0	10.3	99.6	58.5	53.7	88.9	97.8		
MDR	229	58.1	13.1	16.2	88.2	31.9	35.4	80.8	83.2		
Pseudomonas aeruginosa	425	93.2	83.8	80.7	76.9	78.4	71.2	85.2	98.4		
MDR	102	73.5	38.2	36.3	16.7	27.5	13.7	48.0	98.0		
Meropenem-nonsusceptible	98	72.4	44.9	44.9	0.0	36.7	25.5	55.1	99.0		
Ceftazidime-nonsusceptible	82	65.9	24.4	0.0	34.1	8.5	35.4	56.1	98.8		
Piperacillin-tazobactam-nonsusceptible	92	68.5	30.4	18.5	32.6	0.0	31.5	55.4	98.9		
Cefepime-nonsusceptible	69	59.4	0.0	10.1	21.7	7.2	24.6	43.5	100.0		
β-lactam-nonsusceptible	46	45.7	0.0	0.0	0.0	0.0	15.2	34.8	100.0		

Table 2 MIC distribution of ceftolozane-tazobactam for the main pathogens and resistant phenotypes (PACTS, 2012–2017)

			Ceftolozane-tazobactam MIC mg/L (%) ^a													
Organism/organism group	Total	0.015	0.03	0.06	0.12	0.25	0.5	1	2	4	8	16	32	> 32	MIC ₅₀	MIC ₉
Escherichia coli	649		0 (0.0)	7 (1.1)	262 (41.4)	265 (82.3)	67 (92.6)	29 (97.1)	4 (97.7)	5 (98.5)	2 (98.8)	1 (98.9)	1 (99.1)	6 (100.0)	0.25	0.5
ESBL, non-CRE	141			0 (0.0)	5 (3.5)	39 (31.2)	50 (66.7)	29 (87.2)	3 (89.4)	5 (92.9)	2 (94.3)	1 (95.0)	1 (95.7)	6 (100.0)	0.5	4
Klebsiella pneumoniae	287		0 (0.0)	3 (1.0)	35 (13.2)	99 (47.7)	67 (71.1)	27 (80.5)	10 (84.0)	9 (87.1)	6 (89.2)	4 (90.6)	7 (93.0)	20 (100.0)	0.5	16
ESBL, non-CRE	117				0 (0.0)	17 (14.5)	41 (49.6)	21 (67.5)	9 (75.2)	8 (82.1)	6 (87.2)	4 (90.6)	4 (94.0)	7 (100.0)	1	16
Pseudomonas aeruginosa	425			0 (0.0)	2 (0.5)	24 (6.1)	234 (61.2)	95 (83.5)	21 (88.5)	20 (93.2)	7 (94.8)	5 (96.0)	3 (96.7)	14 (100.0)	0.5	4
Ceftazidime-nonsusceptible	82					0 (0.0)	1 (1.2)	19 (24.4)	15 (42.7)	19 (65.9)	6 (73.2)	5 (79.3)	3 (82.9)	14 (100.0)	4	>32
Piperacillin-tazobactam-nonsusceptible	92				0 (0.0)	1 (1.1)	5 (6.5)	23 (31.5)	16 (48.9)	18 (68.5)	7 (76.1)	5 (81.5)	3 (84.8)	14 (100.0)	4	>32
Cefepime-nonsusceptible	69						0 (0.0)	9 (13.0)	13 (31.9)	19 (59.4)	7 (69.6)	5 (76.8)	3 (81.2)	13 (100.0)	4	>32
Meropenem-nonsusceptible	98					0 (0.0)	22 (22.4)	24 (46.9)	11 (58.2)	14 (72.4)	7 (79.6)	4 (83.7)	3 (86.7)	13 (100.0)	2	>32
β-lactam-nonsusceptible	46						0 (0.0)	5 (10.9)	4 (19.6)	12 (45.7)	6 (58.7)	4 (67.4)	3 (73.9)	12 (100.0)	8	>32
MDR	101					0 (0.0)	12 (11.9)	38 (37.6)	56 (55.4)	74 (73.3)	81 (80.2)	85 (84.2)	88 (87.1)	101 (100.0)	2	>32

Table 3 Numbers of isolates by patient age for main organism groups

a EUCAST (2018) susceptible breakpoints shaded

Abbreviations: yo, year old.

Organism group	≤1 yo	2–6 yo	7–12 yo	>13 yo
All isolates	937 (43.9%)	470 (22.0%)	398 (18.6%)	331 (15.5%)
All Enterobacteriaceae	658 (46.4%)	303 (21.4%)	244 (17.2%)	212 (15.0%)
MDR	109 (48.0%)	52 (22.9%)	34 (15.0%)	32 (14.1%)
ESBL, non-CRE	123 (45.4%)	64 (23.6%)	47 (17.3%)	37 (13.7%)
All P. aeruginosa	129 (30.4%)	100 (23.5%)	108 (25.4%)	88 (20.7%)
MDR	31 (30.7%)	19 (18.8%)	25 (24.8%)	26 (25.7%)

• The number of organisms and organism groups by patient age is shown in

 The largest group of patients, ≤1 yo, had the largest number of *Enterobacteriaceae* and *P. aeruginosa* isolates and the largest number of MDR and ESBL, non-CRE

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CONCLUSIONS

- Ceftolozane-tazobactam susceptibility was 89.8% against Enterobacteriaceae, 97.1% against E. coli, and 80.5% against K. pneumoniae paediatric isolates
 - Meropenem and amikacin were the most active antimicrobials against Enterobacteriaceae
- For *P. aeruginosa*, ceftolozane-tazobactam was the most potent β-lactam tested with activity similar to colistin
- Ceftolozane-tazobactam maintained activity against P. aeruginosa isolates nonsusceptible to other β-lactams and MDR isolates
- These data suggest that ceftolozane-tazobactam may be a useful treatment for serious paediatric infections caused by gram-negative pathogens, including *P. aeruginosa*

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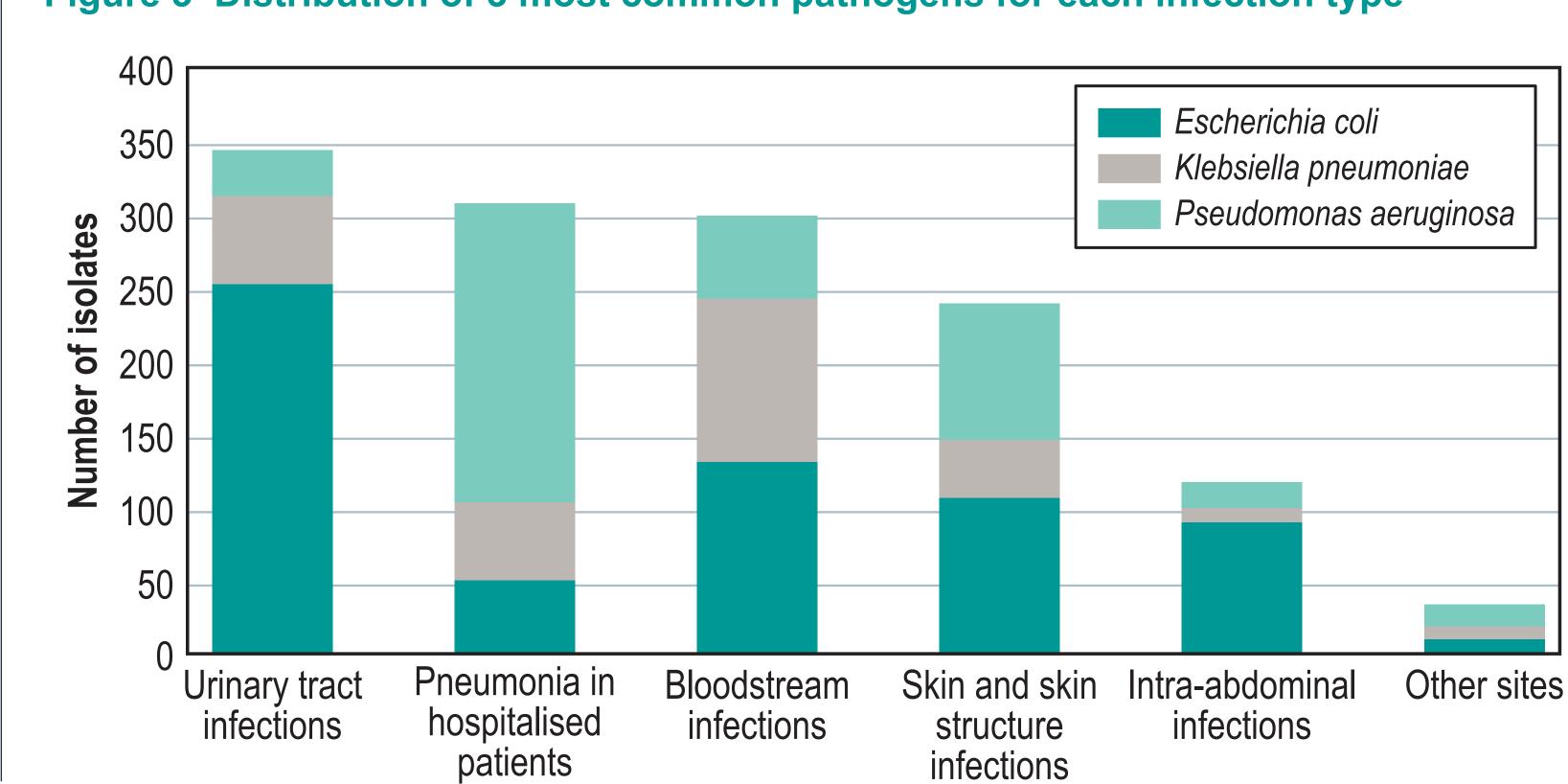
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