

New Chromosomal AmpC β -Lactamase in *Enterobacter cloacae*

Several members of the *Enterobacteriaceae*, including *Enterobacter* spp., are naturally resistant to amoxicillin and cephalosporins. *Enterobacter cloacae* produces chromosomally encoded β -lactamases, also called cephalosporinases (1), and is a serious nosocomial pathogen, the third most prevalent bacterium isolated in intensive care settings (5, 8). We report here the study of a new chromosomal AmpC β -lactamase produced by *E. cloacae* FFUL2En isolated from the blood culture of a patient hospitalized in a medicine ward of Hospital de Santa Maria, Lisbon, Portugal. The antibiogram revealed resistance to aminopenicillins, aztreonam, and broad-spectrum cephalosporins, except imipenem, aminoglycosides, and quinolones. By isoelectrofocusing, the sonicate extracts expressed a pI of 8.68, suggesting the presence of a presumed AmpC enzyme.

A total DNA preparation from *E. cloacae* FFUL2En was used in PCR experiments with two sets of primers, TN5 (5'-CGTTT GTCAGGCACAGTCAAATCCA) and TN4 (5'-TTACTGTAG CGCGTCGAGGATATGG) and the internal primers TN2 (5'-TTCCACTGCGGCTGCCAGT) and TN3 (5'-CGGATGAGG TCACGATAACGCC), designed in accordance with consensus sequences from the *ampC* genes described for *E. cloacae* and available at GenBank. The amplicon with 1,234 bp was cloned into the SmaI site of the pBK-CMV vector (6) with a TOPO TA cloning kit, resulting in the plasmid p2En1. The β -lactam susceptibility pattern of *Escherichia coli* 2En1, harboring the recombinant plasmid p2En1, displayed cefoxitin, cefuroxime, ceftazidime, and piperacillin plus tazobactam MICs of >256 μ g/ml and a cefepime MIC of 0.5 μ g/ml. The MICs of cefotaxime and aztreonam were lower than those for the parental strain (Table 1). The *E. coli* 2En1 transformant showed the same pI as the parental strain (pI 8.68), and the substrate profile of the enzyme EcoFFUL2En was determined with the transformant crude enzymatic extract (7). The V_{max} values indicate that cephalothin, with a V_{max} of 3,000.1 μ M/min, is hydrolyzed more quickly than cefoxitin ($V_{max} = 3.7 \mu$ M/min). Ceftazidime and cefotaxime are not hydrolyzed at detectable levels ($V_{max} = <0.1 \mu$ M/min).

In order to perform the sequencing reactions, the amplicon of 1,234 bp was cloned in the pCR2.1-TOPO vector with a TOPO TA cloning kit, resulting in the plasmid p2En2. The sequence with 382 amino acids has an 86% identity with the AmpC of *E. cloacae* P99 and 98% identity with the plasmid-borne MIR-1 β -lactamase gene product (2).

To search for a possible chromosomal location of the *bla*_{AmpC}

gene, whole-cell DNA of *E. cloacae* FFUL2En was restricted with I-CeuI endonuclease (New England Biolabs), which recognizes a 26-bp sequence in *rm* genes coding for the 23S large-subunit rRNA. After digestion, separation of the resulting fragments was performed on a contour-clamped homogeneous electric field-DRII apparatus, as described previously (3).

The restricted fragments of *E. cloacae* FFUL2En DNA were transferred to a nylon membrane by Southern blotting (9) and were hybridized by using a nonradioactive labeling and detection kit (Roche) with a PCR-obtained probe with primers TN5 and TN2 (see above), consisting of a 576-bp fragment of *bla*_{AmpC} and a 16S rRNA gene probe amplified with universal primers described elsewhere (4). The *bla*_{AmpC} probe hybridized only with the 630-kb fragment of *E. cloacae* FFUL2En. These data indicate the chromosomal location of the *bla*_{AmpC} gene, coding for the AmpC β -lactamase EcoFFUL2En, in *E. cloacae* FFUL2En, which is closely related to the plasmid-borne MIR-1 from *Klebsiella pneumoniae*.

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TABLE 1. MICs of β -lactams for *E. cloacae* FFUL2En clinical isolate, *E. coli* 2En1 harboring recombinant plasmid p2En1, and reference strain *E. coli* TOP10 harboring the pBK-CMV plasmid

β -Lactam	MIC (μ g/ml)		
	<i>E. cloacae</i> FFUL2En	<i>E. coli</i> 2En1	<i>E. coli</i> TOP10(pBK-CMV)
Piperacillin + TZB ^a	>256	>256	3
Cefoxitin	>256	>256	8
Cefuroxime	>256	>256	4
Cefotaxime	>256	8	0.094
Ceftazidime	>256	>256	0.5
Cefepime	0.5	0.38	0.064
Aztreonam	48	6	0.094
Imipenem	0.75	0.38	ND ^b

^a TZB, tazobactam.

^b ND, not determined.