

Development of Test Panel of β -Lactamases Expressed in a Common *Escherichia coli* Host Background for Evaluation of New β -Lactam Antibiotics

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A test panel of 35 different β -lactamases expressed in a common *Escherichia coli* host was created to compare the effect that each β -lactamase had on susceptibility to various β -lactam antibiotics. A comparison of the MICs obtained with this panel generally reflected differences in the substrate profiles of the various β -lactamases examined. In addition, several strains of the panel were subjected to selection with porin-specific bacteriophages to obtain mutants lacking either the OmpC or OmpF porin protein. A mutation in either OmpC or OmpF did change the susceptibilities of certain strains expressing β -lactamase to certain β -lactam antibiotics. However, the loss of a single porin did not predictably alter susceptibility to any given β -lactam drug. This panel of strains producing various β -lactamases was found to be a useful tool for comparing the effects of different β -lactamases and outer membrane permeability upon susceptibility to β -lactam drugs.

The production of β -lactamase is the dominant mechanism for bacterial resistance to β -lactam antibiotics. Both the level of β -lactamase activity and the substrate profile of the enzyme play a role in determining bacterial resistance conferred by the enzyme in gram-negative bacteria (20, 44, 50, 54). In addition, the outer membrane of gram-negative bacteria is known to act as a permeability barrier for various antibiotics, including the β -lactams. The combination of external antibiotic concentration, permeability, and β -lactamase activity determines the periplasmic free drug concentration and, consequently, susceptibility or resistance to the drug (31, 40, 41). The appearance of clinical isolates of the family *Enterobacteriaceae* that have a high level of resistance to β -lactam antibiotics has been shown, in some instances, to be the result of the interaction between β -lactamase production and decreased permeability. For example, the combination of an extended-spectrum β -lactamase and altered outer membrane proteins was responsible for high-level resistance to ceftazidime in a clinical isolate of *Escherichia coli* producing TEM-12 (63).

Because of the importance of β -lactamase and permeability in resistance to β -lactam antibiotics, the present study was undertaken in an attempt to develop a test panel of *E. coli* isolates that would allow an assessment of the effects of various β -lactamases and altered permeability on susceptibility to β -lactam antibiotics. First, a test panel consisting of many different plasmid-mediated β -lactamases expressed in a common host background was constructed. Subsequently, several strains expressing β -lactamase were subjected to selection with porin-specific bacteriophage, resulting in the isolation of mutants lacking either the OmpC or the OmpF porin protein. The resulting panel was then used to (i) compare the effects of these β -lactamases on susceptibilities to various β -lactam antibiotics and (ii) assess the influence of the combination of β -lactamase expression and a change in outer membrane proteins on susceptibility to β -lactam antibiotics.

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MATERIALS AND METHODS

Creation of the test panel. A panel of different plasmid-mediated β -lactamases expressed in a common host background was created in *E. coli* C600N, a nalidixic acid-resistant derivative of *E. coli* C600 (3) (kindly provided by R. V. Goering). The plasmids encoding the β -lactamases used in the study are listed in Table 1. The plasmids were transferred to the recipient by either conjugation or transformation by previously described methods, selecting for resistance to both nalidixic acid and ampicillin (3, 4, 12, 49). The presence of a single β -lactamase of the correct isoelectric point in the strains of the test panel was confirmed by isoelectric focusing (55, 62).

E. coli C600N panel strains constitutively producing high-levels of group 1 β -lactamase (Ecl-C) and producing a high basal level and hyperinducible group 1 β -lactamase (Ecl-Hy) were spontaneous mutants of the wild-type inducible strain (Ecl-I) containing pBP131 with the *ampR ampC* region cloned from *Enterobacter cloacae* 14 (27). An overnight culture of the strain inducibly expressing the group 1 enzyme (Ecl-I) was diluted in fresh Luria-Bertani (LB) broth (Difco Laboratories, Detroit, Mich.) and was grown to the mid- to late log phase. Serial 10-fold dilutions of the cells were made in sterile saline, and 0.1 ml was plated onto LB agar plates containing 8, 16, and 32 μ g of cefotaxime ($2\times$ the MIC). Mutants were selected and tested for constitutive β -lactamase production by induction studies (52, 53).

Selection with bacteriophage. Spontaneous mutants of nine strains of the *E. coli* C600N panel lacking either the OmpC or the OmpF porin protein were selected by using porin-specific bacteriophages. Bacteriophage SS4 is specific for the OmpC of *E. coli* K-12 derivatives, whereas bacteriophage K20 is specific for OmpF (56). Phages were the kind gift of C. A. Schnaitman. Fresh phage lysates were diluted to 10^{-3} in LB broth. A total of 20 μ l of overnight culture and 10 μ l of diluted phage were added to a melted soft agar of nutrient or Trypticase soy agar, and the contents were mixed and poured over the surface of a dry agar plate. Following incubation, spontaneous mutants appear as colonies that grow in the presence of phage. Resistant colonies were subcultured and subjected to reselection with phage. A confluent lawn of bacteria should be obtained from strains that have lost a specific porin. The complete loss of a specific outer membrane protein was confirmed by sodium dodecyl sulfate-polyacrylamide gel electrophoresis analysis in the presence of 8 M urea as described previously (1, 33).

Susceptibility tests. All susceptibility tests were performed by standard procedures (40). Broth microdilution tests were performed with a final inoculum of 10^4 CFU per well. Microdilution trays containing serial twofold dilutions of antibiotic were prepared by using the MIC 2000 apparatus (Dynatech Laboratories, Chantilly, Va.). The MICs of the following antibiotics were determined: ticarcillin and potassium clavulanate (Beecham Laboratories, Bristol, Tenn.),

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TABLE 1. β -Lactamases expressed by strains in the *E. coli* C600N test panel

β -Lactamase	Plasmid	Received from:	Reference
TEM-1	R6K	A. Medeiros	10
TEM-1 (high)	pUC19		64
TEM-2	RP1	A. Medeiros	35
TEM-3	pCFF04	J. Sirot	59
TEM-4	pUD16	G. Paul	43
TEM-5	pCFF14	J. Sirot	60
TEM-7		L. Gutmann	17
TEM-8	HM12G	C. Mabilat	34
TEM-9		A. Harris	61
TEM-10	pJPQ100	J. Quinn	46
TEM-12	pDAW402	D. Weber	63
SHV-1	R1010	A. Medeiros	48
SHV-2	pBP60	B. Wiedemann	25
SHV-3	pUD18	A. Philippon	22
SHV-4	pUD21	A. Philippon	7
SHV-5	pAFF2	L. Gutmann	16
SHV-6	pSLH47	G. Arlet	2
OXA-1	RGN238	G. Jacoby	9
OXA-2	R1818 (R46)	A. Medeiros	8
OXA-3	pIP55	G. Jacoby	35
OXA-4	pMG203	A. Medeiros	38
OXA-5	pMG54	A. Medeiros	38
OXA-6	pUZ8-pMG39	A. Medeiros	38
OXA-7	pMG202	A. Medeiros	38
PSE-1	pMG217	G. Jacoby	37
PSE-2	R140	G. Jacoby	32
PSE-3	Rms149	G. Jacoby	18
PSE-4	pMON705	G. Jacoby	15
SAR-1	pUK657	S. Amyes	47
ROB-1	pMON401	G. Jacoby	30
HMS-1	R997	A. Medeiros	36
CARB-4	pMON1024	G. Jacoby	45
LCR-1	pMK20:Tn1412	G. Jacoby	29, 58
L-1	pMON01	R. Levesque	11
MIR-1	pMLC28Kpn1-8	A. Medeiros	42
ampC	pBP131	G. Korfmann	27

aztreonam and cefepime (Bristol-Meyers-Squibb, Princeton, N.J.), ceftazidime (Glaxo Group Research Ltd., Greenford, England), cefotaxime and ceftiprome (Hoechst-Roussel Pharmaceuticals Inc., Sommerville, N.J.), cefixime, piperacillin, and tazobactam (Lederle Laboratories, Pearl River, N.Y.), ampicillin, cefoperazone, and sulbactam (Pfizer Inc., New York, N.Y.), and amoxicillin (Sigma Chemical Company, St. Louis, Mo.). Ampicillin-sulbactam, amoxicillin-clavulanate, and cefoperazone-sulbactam combinations were tested at 2:1 ratios. Serial twofold dilutions of ticarcillin, each containing 2 μ g of clavulanate per ml, and serial twofold dilutions of piperacillin, each containing 4 μ g of tazobactam per ml, were tested. The MIC was defined as the lowest concentration preventing growth after 18 h of incubation at 35°C. Results for most antibiotics were interpreted by using the criteria approved by the National Committee for Clinical Laboratory Standards (39). For cefoperazone-sulbactam the criteria of Jones et al. (23) were used, for cefepime the criteria of Fuchs et al. (13) were used, and for ceftiprome the criteria of Jones et al. (24) were used. When comparing MIC results, a significant change in MIC was defined as a fourfold increase or decrease in the MIC.

RESULTS

Effect of type of β -lactamase on susceptibility. Broth microdilution MICs were determined for all strains of the *E. coli* C600N test panel (Table 2). In general, the MICs of the β -lactam antibiotics tested followed the reported substrate profiles for the various β -lactamases. The presence of any β -lactamase raised the MIC of ampicillin to ≥ 128 μ g/ml and the MIC of ticarcillin to ≥ 32 μ g/ml. Sulbactam restored susceptibility to ampicillin in strains expressing TEM-3, TEM-4, TEM-8, TEM-9, TEM-10, TEM-12, OXA-2, OXA-3, PSE-1, PSE-3, and ROB-1 β -lactamases. Clavulanate restored susceptibility

to amoxicillin only in the strain expressing PSE-3. Clavulanic acid restored susceptibility to ticarcillin in strains expressing OXA-2, LCR-1, and the inducible group 1 β -lactamase (Ecl-I).

All of the TEM, SHV, and PSE β -lactamases raised the MIC of piperacillin to >128 μ g/ml with the exception of the MICs for strains expressing TEM-5 and PSE-1, which were moderately susceptible to piperacillin (Table 2). Tazobactam did not restore susceptibility to piperacillin in strains expressing TEM-1 or TEM-2; however, the MICs of the combination were 2 to 8 μ g/ml when they were determined for strains expressing TEM-3 through TEM-12. Among strains expressing SHV β -lactamases, tazobactam restored susceptibility to piperacillin only in those strains expressing SHV-2 and SHV-3. All of the strains expressing PSE enzymes were susceptible to the combination. Strains expressing TEM-1, TEM-2, SHV-1, SHV-2, SHV-3, SHV-5, SHV-6, PSE-2, HMS-1, and CARB-4 β -lactamases were resistant to cefoperazone (MICs, 64 to >128 μ g/ml). Sulbactam did not restore susceptibility to cefoperazone in strains expressing TEM-3, SHV-6, HMS-1, or CARB-4.

For strains which were susceptible to the β -lactam- β -lactamase inhibitor combinations, susceptibility was not as great as that of the β -lactamase-negative strain, C600N (Table 2). In general, the MICs of these antibiotics for strains producing a TEM β -lactamase were lower than those for strains producing an SHV enzyme. In addition, MICs for strains producing the extended-spectrum β -lactamases (TEM-3 through TEM-12, SHV-2 through SHV-6) were lower than those for strains producing the parental enzymes TEM-1, TEM-2, or SHV-1. Although the group 1 enzymes are not highly susceptible to the β -lactamase inhibitors, the inhibitors did lower the MICs of the companion drug 2- to 32-fold. However, susceptibility was not restored for any of the combinations with the strains expressing the group 1 enzyme constitutively (Ecl-C) or hyperinducibly (Ecl-Hy) (Table 2).

For the extended-spectrum β -lactam drugs, the presence of TEM-1, TEM-2, and SHV-1 β -lactamases did not significantly affect the MICs of cefotaxime or aztreonam. However, the MICs of ceftazidime and ceftiprome in the presence of TEM-2 and SHV-1 were four- to eightfold higher than the MICs obtained for *E. coli* C600N. For strains expressing an extended-spectrum β -lactamase (TEM-3 through TEM-12, SHV-2 through SHV-6) the MICs of cefotaxime, ceftazidime, aztreonam, cefepime, and ceftiprome were increased up to 4,096-fold. The MICs of cefotaxime were greater than the MICs of ceftazidime for four strains, while the MICs of ceftazidime were greater than the MICs of cefotaxime for eight strains. The MICs of aztreonam were greater than the MICs of either ceftazidime or cefotaxime for the strains expressing the SHV-5 β -lactamase. The extended-spectrum β -lactamases increased the MICs of cefepime and ceftiprome 4- to 128-fold above those obtained in tests with the host strain, C600N.

The presence of any of the group 1 β -lactamases (MIR-1, Ecl-I, Ecl-C, and Ecl-Hy) significantly increased the MICs of cefotaxime, ceftazidime, and aztreonam, regardless of the level of enzyme expression (Table 2). In contrast, only high-level expression of group 1 β -lactamase was sufficient to significantly increase the MICs of cefepime or ceftiprome.

Effect of a change in outer membrane protein on susceptibility. Mutants lacking either *OmpC*, *OmpF*, or both *OmpC* and *OmpF* were selected by using porin-specific phages with strains from the C600N panel expressing TEM-1, TEM-4, TEM-9, SHV-1, SHV-4, MIR-1, Ecl-I, and Ecl-C β -lactamases. The MICs of the various β -lactam drugs determined for these strains are given in Table 2. There was no general trend for the loss of one porin correlating with a change in the MIC.

TABLE 2. MICs of various β -lactam antibiotics for the *E. coli* C600N test panel

Enzyme	Porin	MIC (μ g/ml) ^a														
		AM	AMC	SAM	PIP	P/T	TIC	TIM	CFP	C/S	CFM	CTX ^b	CAZ ^b	ATM	PIME ^b	ROME ^b
None	WT ^c	4	8	4	2	2	8	4	≤0.06	0.125	0.5	≤0.06	0.125	≤0.06	≤0.06	≤0.06
None	OmpC ⁻	4	8	8	2	2	4	4	≤0.06	0.125	0.25	≤0.06	0.25	≤0.06	≤0.06	≤0.06
None	OmpF ⁻	8	8	4	2	2	16	8	0.5	0.5	0.5	≤0.06	0.25	0.125	≤0.06	≤0.06
TEM-1	WT	>128	16	128	>128	32	>128	128	128	8	0.5	≤0.06	0.25	≤0.06	≤0.06	0.125
TEM-1	OmpC ⁻	>128	16	64	>128	32	>128	128	>128	4	0.5	≤0.06	0.5	≤0.06	≤0.06	0.125
TEM-1	OmpF ⁻	>128	32	128	>128	16	>128	128	64	16	0.5	≤0.06	0.5	0.125	0.125	0.25
TEM-1 (high)	WT	>128	32	>128	>128	64	>128	128	128	32	0.25	≤0.06	0.25	0.125	0.25	0.25
TEM-2	WT	>128	64	>128	>128	64	>128	128	>128	16	0.5	≤0.06	0.5	0.125	0.125	0.25
TEM-3	WT	>128	16	4	>128	2	>128	64	32	64	4	32	16	4	0.5	2
TEM-4	WT	>128	16	8	>128	4	>128	128	32	1	16	64	16	16	2	8
TEM-4	OmpC ⁻	>128	16	8	>128	8	>128	64	32	1	8	64	8	8	4	8
TEM-4	OmpF ⁻	>128	16	8	>128	4	>128	128	64	1	16	128	32	16	4	32
TEM-4	OmpC ⁻ OmpF ⁻	>128	16	16	>128	16	>128	128	128	4	32	32	16	16	8	32
TEM-5	WT	>128	16	16	64	8	>128	32	16	2	32	4	64	2	2	2
TEM-7	WT	>128	32	16	>128	8	>128	128	16	1	1	0.125	32	1	2	2
TEM-8	WT	>128	16	8	>128	8	>128	64	16	0.5	8	2	128	8	2	2
TEM-9	WT	>128	16	8	>128	8	>128	128	16	0.5	8	2	>128	8	2	4
TEM-9	OmpC ⁻	>128	16	8	>128	8	>128	32	16	1	8	2	>128	8	4	8
TEM-9	OmpF ⁻	>128	16	8	>128	4	>128	128	32	0.5	16	8	>128	32	16	16
TEM-9	OmpC ⁻ OmpF ⁻	>128	16	8	>128	4	>128	128	32	2	16	4	>128	16	8	16
TEM-10	WT	>128	16	8	>128	8	>128	128	8	1	16	1	>128	128	2	2
TEM-12	WT	>128	32	8	>128	8	>128	128	16	1	0.5	≤0.06	16	0.5	2	4
SHV-1	WT	>128	32	>128	>128	128	>128	128	>128	16	0.25	≤0.06	1	≤0.06	0.125	0.25
SHV-1	OmpC ⁻	>128	32	>128	>128	128	>128	128	>128	32	1	≤0.06	1	1	0.5	1
SHV-1	OmpF ⁻	>128	>128	>128	>128	128	>128	128	128	32	1	≤0.06	1	0.25	0.25	1
SHV-1	OmpC ⁻ OmpF ⁻	>128	32	>128	>128	32	>128	128	>128	8	0.25	≤0.06	1	≤0.06	≤0.06	0.125
SHV-2	WT	>128	16	16	>128	16	>128	128	64	8	2	16	8	4	4	8
SHV-3	WT	>128	16	32	>128	16	>128	32	128	4	4	64	4	4	4	4
SHV-4	WT	>128	16	16	>128	32	>128	128	32	8	>128	32	>128	>128	4	4
SHV-4	OmpC ⁻	>128	8	8	>128	16	>128	128	8	2	64	32	>128	>128	0.5	1
SHV-4	OmpF ⁻	>128	8	8	>128	32	>128	64	64	4	128	64	>128	128	1	2
SHV-5	WT	>128	64	64	>128	32	>128	128	128	16	0.5	0.25	0.25	1	0.25	1
SHV-6	WT	>128	>128	>128	>128	>128	128	128	>128	64	4	1	32	0.5	2	8
OXA-1	WT	>128	64	64	>128	64	>128	128	1	1	0.5	0.5	0.25	0.125	2	2
OXA-2	WT	>128	16	8	32	2	>128	8	4	0.5	0.5	≤0.06	1	≤0.06	≤0.06	≤0.06
OXA-3	WT	>128	32	8	32	4	>128	64	8	1	1	≤0.06	0.5	0.125	≤0.06	0.125
OXA-4	WT	>128	64	32	128	32	>128	128	0.5	0.5	0.25	≤0.06	0.125	≤0.06	0.5	1
OXA-5	WT	128	16	16	4	1	>128	32	1	1	0.5	≤0.06	0.125	≤0.06	≤0.06	≤0.06
OXA-6	WT	>128	16	32	32	8	>128	128	8	8	≤0.06	≤0.06	0.25	0.5	≤0.06	≤0.06
OXA-7	WT	>128	64	64	128	32	>128	128	32	16	0.5	≤0.06	0.125	1	0.125	0.5
PSE-1	WT	>128	32	8	32	1	>128	128	8	0.5	0.25	≤0.06	0.5	≤0.06	≤0.06	≤0.06
PSE-2	WT	>128	32	32	>128	1	>128	128	>128	1	0.5	≤0.06	0.25	≤0.06	≤0.06	≤0.06
PSE-3	WT	>128	8	8	>128	4	>128	64	8	0.5	0.5	≤0.06	0.25	≤0.06	≤0.06	≤0.06
PSE-4	WT	>128	32	64	>128	16	>128	128	32	8	0.25	≤0.06	0.25	≤0.06	0.125	≤0.06
SAR-1	WT	>128	16	32	>128	8	>128	64	16	2	0.5	≤0.06	0.125	≤0.06	≤0.06	≤0.06
ROB-1	WT	>128	16	8	>128	2	>128	128	32	1	0.5	≤0.06	8	0.125	0.25	0.5
HMS-1	WT	>128	32	128	>128	32	>128	128	>128	64	0.5	≤0.06	0.25	≤0.06	≤0.06	≤0.06
CARB-4	WT	>128	32	64	>128	64	>128	128	>128	32	0.25	≤0.06	0.25	≤0.06	0.25	≤0.06
LCR-1	WT	>128	32	32	8	2	64	16	2	1	0.25	0.125	0.25	≤0.06	0.125	0.125
L-1	WT	>128	>128	128	128	>128	>128	128	>128	>128	8	16	64	≤0.06	≤0.06	1
MIR-1	WT	>128	64	64	64	32	>128	128	>128	8	>128	128	64	64	0.25	0.5
MIR-1	OmpC ⁻	>128	128	64	64	32	>128	128	128	16	>128	>128	64	64	1	2
MIR-1	OmpF ⁻	>128	128	32	32	16	>128	128	64	8	>128	32	64	32	0.125	0.25
(Ecl-I)	WT	>128	128	64	16	4	32	16	4	1	64	4	2	0.5	≤0.06	≤0.06
(Ecl-I)	OmpC ⁻	>128	128	64	32	16	128	64	8	4	128	8	16	2	≤0.06	≤0.06
(Ecl-I)	OmpF ⁻	>128	128	64	16	2	64	32	8	4	>128	8	4	1	≤0.06	0.125
(Ecl-I)	OmpC ⁻ OmpF ⁻	>128	128	64	8	4	32	8	2	1	64	8	1	0.5	≤0.06	≤0.06
(Ecl-C)	WT	>128	128	64	>128	64	>128	128	>128	32	>128	128	>128	8	1	8
(Ecl-C)	OmpC ⁻	>128	64	128	128	64	128	128	>128	16	>128	64	128	16	2	64
(Ecl-C)	OmpF ⁻	>128	>128	64	>128	32	>128	128	>128	32	>128	32	>128	16	4	64
(Ecl-C)	OmpC ⁻ OmpF ⁻	>128	>128	64	128	32	128	128	128	32	>128	128	>128	16	2	64
(Ecl-Hy)	WT	>128	64	32	128	64	128	128	128	16	>128	128	128	32	1	16

^a MICs were determined in broth microdilution tests with 10⁴ CFU per well (39). AM, ampicillin; AMC, amoxicillin-clavulanate; SAM, ampicillin-sulbactam; PIP, piperacillin; P/T, piperacillin-tazobactam; TIC, ticarcillin; TIM, ticarcillin-clavulanate; CFP, cefoperazone; C/S, cefoperazone-sulbactam; CFM, cefixime; CTX, ceftotaxime; CAZ, ceftazidime; ATM, aztreonam; PIME, cefepime; ROME, ceftiprome.

^b Some data were published previously (51).

^c WT, wild-type porins (both OmpC and OmpF porin proteins were present).

For example, for the strain expressing TEM-9, the loss of OmpC resulted in a fourfold decrease in the MIC of ticarcillin-clavulanate, while the loss of OmpF resulted in a fourfold or greater increase in the MICs of cefotaxime, aztreonam, cefepime, and ceftiofime (Table 2). In general, penicillins were less affected by changes in outer membrane proteins, whereas ceftiofime was most often affected.

DISCUSSION

Previous studies have used a test panel of β -lactamase-producing *E. coli* to study the activities of various enzymes against β -lactam antibiotics (19, 21). However, those studies included a limited number of types of β -lactamase, and the same host strain was not used throughout all analyses. The test panel presented in the current study was more extensive in its construction in that it included representatives of each of the major groups of β -lactamases, i.e., cephalosporinases, broad-spectrum enzymes, extended-spectrum β -lactamases, enzymes hydrolyzing carbenicillin or cloxacillin, and metallo- β -lactamases. In addition, the panel included mutants of the same strain that differed only in their outer membrane proteins or in the expression of the same enzyme, i.e., the group 1 β -lactamase and the TEM-1 β -lactamase. Therefore, with this panel it was possible to assess the influence of the type of β -lactamase, the level of expression for certain enzymes, and permeability, alone and in combination, upon susceptibility to β -lactam antibiotics.

The results of our study with the porin mutants supported and expanded upon those of Jacoby and Sutton (21) and Jacoby and Carreras (19). Those earlier studies also showed that variations in outer membrane porins had relatively little effect on the level of β -lactamase-mediated resistance. These results may at first seem surprising in light of the high level of resistance that a combination of a change in outer membrane proteins and β -lactamase production can contribute to clinical isolates (50, 54, 63). The lack of any consistent effect seen in these test panels may be due, in part, to the fact that these mutant strains were selected under phage pressure and may therefore behave differently from clinical isolates selected under antibiotic pressure. Also, since the same porin mutant may arise from mutations in either structural or control genes, this may explain the lack of a consistent effect as well.

It should be noted that this and the previous test panels (19, 21) consisted of laboratory strains of *E. coli*. The introduction of β -lactamases into these strains and the selection of mutants altered in enzyme expression or outer membrane permeability may not accurately predict the actual level of resistance which may be seen when the same change occurs in a clinical isolate of *E. coli* because other factors influencing susceptibility to β -lactam antibiotics may differ in clinical isolates. Nevertheless, such test panels can be very useful in assessing whether a specific factor, taken alone or in combination with others, will influence susceptibility to a given β -lactam antibiotic. However, since the plasmids encoding the various β -lactamases were not the same, differences in plasmid copy number and promoter efficiency could be responsible for some of the differences in susceptibility observed across the various enzymes.

A number of interesting observations were made with the test panel used in the current investigation. First, β -lactamase inhibitors did not always restore susceptibility to their companion drugs in tests with strains producing enzymes highly susceptible to the inhibitors. This supports reports of clinical isolates resistant to inhibitor-drug combinations because of the

hyperproduction of inhibitor-susceptible enzymes (52, 54). Second, the diminished impacts of extended-spectrum β -lactamases on the activities of cefepime and ceftiofime in comparison with their effects on older expanded-spectrum β -lactam antibiotics were apparent in tests with the panel. Whether this predicts the clinical success of the new drugs for the treatment of infections caused by strains producing extended-spectrum β -lactamases awaits human trials. However, it should be noted that the acquisition of certain extended-spectrum β -lactamases did diminish the activities of cefepime and ceftiofime.

There were also several interesting observations that were made with the porin mutants examined in the present study. First, the activities of both cefepime and ceftiofime were diminished by the loss of OmpC, OmpF, or both porins in strains constitutively producing the group 1 β -lactamase. These drugs are reported to be more active against hyperproducers of this β -lactamase than are other cephalosporins in part because they diffuse into the bacterial cell at a faster rate (14, 26, 28, 41, 51, 53, 57). Therefore, it was not surprising to see an increase in the level of resistance to these compounds caused by a decrease in permeability in strains expressing high levels of group 1 β -lactamase. However, this effect was not seen in the strain producing the MIR-1 enzyme. The difference between the strains producing MIR-1 and the group 1 β -lactamase constitutively may lie either in differences between the enzymes themselves, the lack of *ampD* in the latter strain, or both factors. Overall, results of the study with the outer membrane protein mutants suggest that while a loss of either OmpC or OmpF did change the susceptibilities of certain strains expressing β -lactamase to specific β -lactam antibiotics, the loss of a single porin did not predictably alter the susceptibility of a strain containing β -lactamase to any given β -lactam drug.

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