

## Pharmacodynamic Evaluation of Factors Associated with the Development of Bacterial Resistance in Acutely Ill Patients during Therapy

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**The selection of bacterial resistance was examined in relationship to antibiotic pharmacokinetics (PK) and organism MICs in the patients from four nosocomial lower respiratory tract infection clinical trials. The evaluable database included 107 acutely ill patients, 128 pathogens, and five antimicrobial regimens. Antimicrobial pharmacokinetics were characterized by using serum concentrations, and culture and sensitivity tests were performed daily on tracheal aspirates to examine resistance. Pharmacodynamic (PD) models were developed to identify factors associated with the probability of developing bacterial resistance. Overall, in 32 of 128 (25%) initially susceptible cases resistance developed during therapy. An initial univariate screen and a classification and regression tree analysis identified the ratio of the area under the concentration-time curve from 0 to 24 h to the MIC ( $AUC_{0-24}/MIC$ ) as a significant predictor of the development of resistance ( $P < 0.001$ ). The final PK/PD model, a variant of the Hill equation, demonstrated that the probability of developing resistance during therapy increased significantly when antimicrobial exposure was at an  $AUC_{0-24}/MIC$  ratio of less than 100. This relationship was observed across all treatments and within all organism groupings, with the exception of  $\beta$ -lactamase-producing gram-negative organisms (consistent with type I  $\beta$ -lactamase producers) treated with  $\beta$ -lactam monotherapy. Combination therapy resulted in much lower rates of resistance than monotherapy, probably because all of the combination regimens examined had an  $AUC_{0-24}/MIC$  ratio in excess of 100. In summary, the selection of antimicrobial resistance appears to be strongly associated with suboptimal antimicrobial exposure, defined as an  $AUC_{0-24}/MIC$  ratio of less than 100.**

Increasing bacterial resistance, and the subsequent burden to society in terms of morbidity, mortality, and increased health care expenditures, necessitates innovative approaches to the use of antimicrobial therapy (2, 15). Considering the dearth of available innovative approaches, attention to the appropriate utilization of antimicrobials is becoming increasingly important, particularly as there are fewer new antimicrobial drugs in development (4). The elucidation of relationships between pharmacodynamic parameters and organism persistence or resistance during therapy would facilitate the design of more effective dosing regimens. Unfortunately, there have been relatively few pharmacodynamic examinations of the relationship between antibiotic dosing and resistance in patients. Most of the available data comes from animal or in vitro models integrating pharmacokinetics and pharmacodynamics, where there are actually many definitive studies and considerable amounts of data.

Reports on the relationship between antimicrobial pharmacodynamic parameters and clinical and microbiological outcomes identify the percent time above the MIC (time-dependent killing) as the parameter predictive of response to  $\beta$ -lactam antimicrobials (8, 17, 27) and the ratio of the peak concentration to the MIC or area under the concentration-time curve (AUC) to the MIC (concentration-dependent kill-

ing) as the predictor for response to aminoglycosides and fluoroquinolones (17, 19). Recently, the AUC-to-MIC ratio, which incorporates both concentration intensity and exposure over time, has been advocated for use as the parameter for prediction of bacteriologic response (14, 16, 18, 26).

We have previously described the pharmacodynamic parameter area under the inhibitory time curve (AUIC) and its relationship to bacteriologic eradication and clinical outcome in patients with nosocomial pneumonia (14, 16, 23, 24). Antimicrobial resistance developed to some extent in each of these trials, but in the individual trials the numbers were too small to clearly define a mathematical relationship. The purpose of this pharmacokinetic/pharmacodynamic analysis was to determine the relationship of antimicrobial exposure, expressed as the AUC/MIC ratio, antibacterial activity, and other covariates to the development of bacterial resistance in the entire patient population treated in these trials.

### MATERIALS AND METHODS

All patients treated in four antimicrobial clinical trials conducted at the Millard Fillmore Hospital between 1984 and 1991 were reviewed (5, 7, 12, 22, 24). The four data sets included a total of 143 acutely ill patients, virtually all of whom were treated for lower respiratory tract infection (LRTI). The four trials included (i) an open-label study of cefmenoxime therapy, 1 to 2 g every 4 or 6 h; (ii) an open-label trial of intravenous ciprofloxacin, 200 to 300 mg every 12 h; (iii) a multicenter, double-blind, randomized trial comparing intravenous ciprofloxacin, 400 mg every 8 h, with intravenous imipenem, 1,000 mg every 8 h; and (iv) an open-label, randomized, antimicrobial exposure (target AUIC of  $250 \text{ SIT}^{-1} \times 24 \text{ h}$  [inverse serum inhibitory titer integrated from 0 to 24 h]) controlled study of intravenous ciprofloxacin (400 mg every 8 or 12 h) versus intravenous ceftazidime (1 to 2 g every 8 or 12 h). In this study, if the dosage needed to provide a target AUC/MIC ratio of 350 exceeded 1,200 mg or 6 g per day for ciprofloxacin,

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TABLE 1. Baseline characteristics by clinical trial

Characteristic or disease condition	% of cases in indicated antibiotic treatment group				
	All studies (n = 107)	Cefmenoxime (n = 26)	Ciprofloxacin (n = 39)	Imipenem vs ciprofloxacin (n = 8)	Ceftazidime vs ciprofloxacin (n = 34)
Ventilator dependent	72.9	76.9	59.0	75.0	85.3
COPD	32.7	42.3	23.1	25.0	38.2
Major surgery	36.4	23.1	35.9	37.5	44.1
Diabetes mellitus	25.3	11.5	25.6	0	41.2
Steroids	25.2	26.9	23.1	12.5	29.4
Malignancy or neoplasm	14.0	23.1	12.8	0	8.8
Chemotherapy or radiation therapy	3.7	11.5	2.6	0	0
Microbiologic cure	67.6	53.8	64.1	62.5	84.8
Clinical cure	68.3	65.4	61.5	100	72.7
Bacterial eradication	67.0	46.1	64.1	87.5	82.3
Prior antibiotics	63.2	84.6	79.5	0	44.1
Serum albumin					
≥3.9 g/dl	1.9	0	0	28.6	0
3.0–3.8 g/dl	22.4	19.2	23.1	14.3	29.0
1.9–2.9 g/dl	72.0	76.9	76.9	42.9	64.5
<1.9 g/dl	3.7	3.8	0	14.3	6.5

cin or ceftazidime, respectively, then piperacillin was added to ciprofloxacin and tobramycin was added to ceftazidime to achieve the targeted AUC.

All patients with LRTI from the above trials were eligible for evaluation. Exclusion criteria were the following: an infection other than an aerobic bacterial pneumonia (anaerobic infection, lung abscess, fungal infection, atypical pathogens); less than 48 h of antimicrobial therapy; inability to isolate a bacterial pathogen; lack of MIC data; and the absence of pharmacokinetic data.

Initial bacteriologic studies on tracheal aspirates were performed for the four clinical trials by the Millard Fillmore Hospital Clinical Laboratory, Department of Pathology. Standard methods for pathogen identification and susceptibility testing using microdilution techniques were employed as previously described (12, 14, 16, 22–24). Culture and sensitivity testing was performed daily, in most cases, throughout the course of therapy and during the follow-up period, for determination of microbiologic end points. Criteria for clinical and microbiologic cure were similar in all studies. The study end points were time to eradication and microbiologic cure. These study end points allowed for the determination of the extent of and the time to the development of bacterial resistance. All isolates considered as bacterial pathogens in the original studies were included in the analysis of the development of resistance. Tracheal-aspirate culture and sensitivity data were reviewed for any significant changes in MICs during therapy. The development of bacterial resistance was defined as the isolation of a bacterial strain, initially found to be susceptible to the treatment regimen, which tested resistant during therapy and/or follow-up. The National Committee for Clinical Laboratory Standards (21) document of MIC interpretive standards was utilized for defining the antimicrobial resistance breakpoints. The cefmenoxime resistance breakpoint was defined by the cefmenoxime study protocol as a MIC of  $\geq 25$   $\mu\text{g}/\text{ml}$ . Time to bacterial resistance was the number of days from the isolation of an initially susceptible organism to the day when the MIC reached the resistance breakpoint.

As there were a large number of different bacterial strains in these patients, the bacterial pathogens were categorized into four groups based upon similar initial-susceptibility characteristics. Group 1 contained only *Pseudomonas* spp. Group 2 contained gram-negative organisms, typically resistant to narrow-spectrum cephalosporins (cefazolin or cephalothin) and whose characteristics were consistent with those of type I  $\beta$ -lactamase producers, such as *Enterobacter cloacae*, *Enterobacter aerogenes*, *Serratia marcescens*, *Citrobacter* spp., *Morganella morganii*, and *Proteus vulgaris*. Group 3 contained other gram-negative rods typically susceptible to cephalosporins, such as *Escherichia coli*, *Klebsiella pneumoniae*, and *Proteus mirabilis*. Group 4 contained the remainder of the diverse organisms, i.e., *Haemophilus influenzae*, *Staphylococcus aureus*, and *Streptococcus pneumoniae*.

Patient-specific pharmacokinetic and pharmacodynamic parameters were determined from serial blood samples and culture MICs. Ciprofloxacin, ceftazidime, cefmenoxime, and piperacillin serum drug concentrations were determined by high-performance liquid chromatography as previously described (5, 7, 22, 23, 24). Specific drug AUC values were obtained by fitting pharmacokinetic models to drug concentration data and integrating the drug concentration-versus-time curves over time. The pharmacodynamic parameter  $\text{AUC}_{0-24}/\text{MIC}$ , where  $\text{AUC}_{0-24}$  is the AUC from 0 to 24 h, was calculated from each subject's specific AUC and organism-specific MIC. In the first three studies, dosing adjustments were infrequent and the initial  $\text{AUC}_{0-24}/\text{MIC}$  ratio was considered to be an adequate estimate of average antimicrobial exposure for the duration of therapy for these cases. In the final study, dose adjustments were made based on initial and subsequent serum concentrations by utilizing an adaptive feedback control

algorithm to maintain targeted antimicrobial exposure ( $\text{AUC}_{0-24}/\text{MIC}$ ). In addition to serum drug concentrations, SITs were also obtained (5, 7). For these cases an average daily  $\text{AUC}_{0-24}/\text{MIC}$  measure was determined by comodeling patient-specific pharmacokinetic and pharmacodynamic parameters with ADAPT II software (9, 10).

**Statistical analysis.** For univariate analysis, the effect of categorical data on the likelihood of the development of resistance was evaluated by the chi-square test and Fisher's exact test where appropriate. Continuous data, such as the patient baseline characteristics of age, weight, and time to resistance, were compared among groups by using Kruskal-Wallis analysis of variance. A *P* value of less than 0.05 was indicative of statistical significance. The data set was also subjected to classification and regression tree (CART) analysis with SYSTAT software (Systat, Inc., Evanston, Ill.) to identify possible significant interacting factors impacting the dependent variable, the development of resistance. Initial analysis by CART suggested that the  $\text{AUC}_{0-24}/\text{MIC}$  ratio and several interacting factors such as prior antimicrobial therapy and the specific combinations of antimicrobial agent and organism (e.g., cefmenoxime therapy and *Pseudomonas* spp. and  $\beta$ -lactam monotherapy and group 2 organisms) were important. The final pharmacodynamic model, a variant of the Hill equation, described a relationship between antimicrobial exposure, expressed as the  $\text{AUC}_{0-24}/\text{MIC}$  ratio, and the development of resistance for the majority of cases. Models were fit by the maximum-likelihood approach available in ADAPT II. Weighting was by the fitted-inverse observation variance, and model discrimination was accomplished by using Akaike's information criterion (1). Kaplan-Meier survival curves were constructed to assess the probability of developing resistance beginning with the initiation of therapy.

## RESULTS

A total of 143 patients were enrolled in the four clinical trials. Study inclusion criteria were met by 107 patients, with 128 organisms being evaluable. The mean age of the patients  $\pm$  standard deviation was  $68.6 \pm 11.7$  years. There were 64 males (60%) and 43 females (40%), with a mean weight of  $69.4 \pm 16.4$  kilograms. The mean duration of therapy was  $10.7 \pm 3.5$  days, with a range of 3 to 31 days. Patient baseline underlying disease states and case characteristics are listed in Table 1. Thirty-six patients were excluded from the evaluation; details concerning them are listed in Table 2. Serum concentrations were obtained for only ciprofloxacin in the third trial (ciprofloxacin versus imipenem); therefore, for trial 3 only the patients receiving ciprofloxacin were included in this analysis. Of the 128 organisms obtained from 107 patients, a single pathogen was evaluated for 90 patients, two pathogens were evaluated for 13 patients, and three pathogens were evaluated for 4 patients. Table 3 lists all organisms. Baseline MICs for selected pathogens are shown. Resistance developed in 32 (25%) of the evaluable cases. Three additional organisms, *K. pneumo-*

TABLE 2. Patients excluded from the analysis

Study	No. of patients		Reason for patient exclusion				
	In study	Excluded	≤48 H of therapy	No organism	NO PK or PD <sup>a</sup>	Resistant at baseline	Missing case report
Cefmenoxime	30	4	1			3	
Ciprofloxacin	50	11	2	3	5		1
Imipenem vs ciprofloxacin	14	6		3		3	
Ceftazidime vs ciprofloxacin	49	15	5	5	2	3	

<sup>a</sup> PK, pharmacokinetic data; PD, pharmacodynamic data.

*nia* and two strains of *E. cloacae*, were isolated during therapy or during the follow-up period, and were found to be resistant. These organisms were not isolated at baseline; therefore, they were not included in the analysis.

The rates of resistance development as determined for organism groupings are listed in Table 4. The greatest frequency of selected resistance was observed for *Pseudomonas* (group 1 organisms) (46.1%), followed by group 2 organisms (27%) and group 3 organisms (10%). Resistance was not observed in the diverse group of remaining organisms (group 4). When resistance was evaluated by treatment, the greatest rate of resistance was observed for cefmenoxime (42.9%), followed by ciprofloxacin (27.6%), ceftazidime (20%), and the ceftazidime-tobramycin combination (9.1%). Resistance was not observed in the ciprofloxacin-piperacillin combination treatment arm. For monotherapy the rate of selected resistance development was 30.7% (31 of 101 cases), while resistance developed in only 3.7% (1 of 27) of the cases of combination therapy. The rates of resistance development by treatment groups, monotherapy and combination therapy, and by organisms are listed in Table 4.

An initial univariate screen of patient, organism, and antimicrobial factors and their relationship to the development of resistance revealed that age, sex, weight, ventilator status, surgery, chronic obstructive pulmonary disease (COPD), diabetes mellitus, steroid use, malignancy, chemotherapy and/or radiation therapy, ciprofloxacin therapy, ceftazidime therapy, and the presence of group 2 gram-negative rods were not significant as predictors of resistance. However, several factors, including the  $AUC_{0-24}/MIC$  ratio, the presence of group 1 or

group 3 gram-negative rods, cefmenoxime therapy, group 2 organisms treated with  $\beta$ -lactam monotherapy, and previous antimicrobial therapy, were determined to be significant by univariate analysis (Table 5).

The median  $AUC_{0-24}/MIC$  values by organism and treatment groupings are presented in Table 6. The designation of susceptible or resistant indicates the antimicrobial exposure and applies to those organisms that either remained susceptible or became resistant, respectively, during treatment. Although the parameter  $AUC_{0-24}/MIC$  has great variability, there is an apparent trend of emergent resistance at lower levels of antimicrobial exposure.

CART analysis identified four factors as significant:  $AUC_{0-24}/MIC$  ratio, cefmenoxime treatment, and organisms of group 1 and group 2. Treatment and organism interactions, such as *Pseudomonas* treated with cefmenoxime and  $\beta$ -lactamase-producing organisms treated with cefmenoxime and/or ceftazidime, were scrutinized for possible significance. Further inspection of these findings and analysis with pharmacodynamic models revealed that *Pseudomonas* spp. and all other organisms, with the exception of  $\beta$ -lactamase-producing organisms (group 2 organisms) treated with  $\beta$ -lactam monotherapy (cefmenoxime or ceftazidime), exhibited an inverse relationship between the probability of developing resistance and the  $AUC_{0-24}/MIC$  ratio.

The final pharmacodynamic model describing this relationship is represented by the equation  $\%P = [P_0 - (P_0 - P_\infty) \cdot AUIC^H / (AUIC_m^H + AUIC^H)] \cdot (1 - R_2) + P_2 \cdot R_2$ , where  $P_0$  is the asymptotic maximum percent probability of resistance as the  $AUC_{0-24}/MIC$  ratio goes to 0;  $P_\infty$  is the asymptotic minimum percent probability of resistance as the  $AUC_{0-24}/MIC$  ratio goes to infinity;  $R_2$  is an indicator, either 0 or 1, of group 2 organisms ( $\beta$ -lactamase-producing organisms treated with  $\beta$ -lactam monotherapy);  $P_2$  is the percent resistance for cases when  $R_2$  is 1;  $AUIC_m$  is the  $AUC_{0-24}/MIC$  ratio at which  $\%P = 0.5 \cdot (P_0 + P_\infty)$ ; and  $H$  is Hill's constant, which reflects the degree of sigmoidicity. Parameters fitted by this model were  $\%P_0 = 82.6\%$ ,  $\%P_{\min} = 9.2\%$ ,  $\%P_2 = 64.3\%$ ,  $AUC_{0-24}/MIC$  ( $AUIC_m$ ) = 100, and  $H = 40$  (fixed). A log-linear regression approach with weighting was utilized to determine the line of best fit for the group 2 organisms treated with  $\beta$ -lactam therapy. The equation describing the line is  $y = 1.409 - 0.2548 \cdot \log_{10} AUC_{0-24}/MIC$ . The final model fitted the data extremely well. The results of the model goodness-of-fit analysis are presented in Table 7. As described by this model, an

TABLE 3. Median MICs for study organisms at baseline and at the end point (resistance)

Organism (n) <sup>a</sup>	Median MIC ( $\mu\text{g/ml}$ ) of <sup>b</sup> :					
	Ceftazidime		Cefmenoxime		Ciprofloxacin	
	Pre TX	Resistant	Pre TX	Resistant	Pre TX	Resistant
<i>E. aerogenes</i> (8)	8.1 (2)	>32 (1)	1.6 (2)	37.5 (2)	0.02 (4)	4 (1)
<i>E. cloacae</i> (11)	8.2 (2)	NR	1.6 (3)	50 (3)	0.02 (6)	NR
<i>E. coli</i> (14)	0.2 (4)	NR	0.5 (4)	NR	0.01 (6)	NR
<i>H. influenzae</i> (6)	0.4 (2)	NR	ND	NR	0.01 (4)	NR
<i>K. pneumoniae</i> (17)	2.0 (3)	64 (1)	1.0 (3)	NR	0.1 (11)	8 (3)
<i>P. aeruginosa</i> (37)	6 (4)	>32 (1)	12.5 (9) <sup>c</sup>	50 (5)	0.5 (24)	5 (12)
<i>P. mirabilis</i> (6)	0.1 (1)	NR	3.1 (2)	NR	0.06 (4)	NR
<i>S. aureus</i> (5)	8 (1)	NR	ND	NR	0.4 (4)	NR
<i>S. marcescens</i> (11)	0.7 (3)	32 (1)	0.4 (3)	25 (1)	0.1 (5)	NR

<sup>a</sup> Other organisms (number of isolates): *Acinetobacter calcoaceticus* (1); *Citrobacter freundii* (1); *Citrobacter* sp. (1); *Klebsiella oxytoca* (3); *M. morgani* (1); *Pseudomonas fluorescens* (2); *P. vulgaris* (2); one was resistant [MIC = 50]; *S. pneumoniae* (1); *Serratia rubrifaciens* (1).

<sup>b</sup> Pre TX, baseline value; Resistant, end point value; NR, no resistance; ND, not done. Values in parentheses are numbers of isolates.

<sup>c</sup> Range, 3.1 to 12.5  $\mu\text{g/ml}$ .

TABLE 4. Selected resistance rates by organism grouping and treatment

Group <sup>a</sup>	No. of isolates	% R <sup>b</sup> (n)	Rate of resistance development <sup>c</sup> (%) for indicated treatment				
			CIP	CEFMNX	CAZ	CIP/PIP	CAZ/TOB
1	39	46.1 (18)	66.7 (12/18)	55.5 (5/9)	0 (0/3)	0 (0/7)	50 (1/2)
2	37	27 (10)	6.7 (1/15)	70 (7/10)	50 (2/4)	0 (0/3)	0 (0/5)
3	40	10 (4)	17.6 (3/17)	0 (0/9)	20 (1/5)	0 (0/5)	0 (0/4)
4	12	0 (0)	0 (0/8)	ND	0 (0/3)	0 (0/1)	ND
All organisms	131	25 (32)	27.6 (16/58)	42.9 (12/28)	20 (3/15)	0 (0/16)	9.1 (1/11)

<sup>a</sup> For a description of organisms in each group, see Materials and Methods.

<sup>b</sup> R, overall rate of resistance development.

<sup>c</sup> CIP, ciprofloxacin; CEFMNX, cefmenoxime; CAZ, ceftazidime; CIP/PIP, ciprofloxacin plus piperacillin; CAZ/TOB, ceftazidime plus tobramycin; ND, not done. The numbers in parentheses are numbers of resistant isolates/total numbers of isolates.

inverse-effect relationship, which applies to all organisms and treatments with the exception of  $\beta$ -lactam monotherapy for group 2 organisms (consistent with type I  $\beta$ -lactamase-producing gram-negative organisms), exists. The observed and modelled data are graphically presented in Fig. 1. As shown in the figure the solid line represents the modelled response surface for all cases, classified as either susceptible or resistant, within the data set for which the  $AUC_{0-24}/MIC$  inverse-effect relationship applied. The dashed line represents the line of best fit for the group 2 organisms treated with  $\beta$ -lactam monotherapy. Observed cases identified by symbols and numbers are plotted as single points within an  $AUC_{0-24}/MIC$  ratio category at the median value for that category.

For those cases in which the model fits applied, *Pseudomonas aeruginosa* treated with ciprofloxacin represented a large number of the fitted cases. The observed percent resistance for ciprofloxacin monotherapy for pseudomonas organisms was 66.7%. This extremely high incidence of resistance was associated with the level of antimicrobial exposure. For example, the observed percent resistance for pseudomonas organisms treated with ciprofloxacin monotherapy, determined by utilizing the fitted  $AUC_{0-24}/MIC$  breakpoint, was 100% (10 of 10 cases) when the  $AUC_{0-24}/MIC$  ratio was <100 and 25% (2 of 8 cases when it was  $\geq 100$ ). In the case of group 3 organisms (other gram-negative rods), including all treatments, the observed percent resistance was 100% (2 of 2 cases) below the breakpoint, and 5.3% (2 of 38 cases) above the breakpoint. The observed data points of  $\beta$ -lactam monotherapy for group 2 organisms are clearly not reflective of the modelled response surface. The high percentage of resistance (>60%) within this group could not be explained by the pharmacodynamic measure of antimicrobial exposure,  $AUC_{0-24}/MIC$ , as resistance occurred throughout a large range of  $AUC_{0-24}/MIC$  ratios (217 to 14,190).

The median time to the observation of selected bacterial resistance was 6 days. The median time to resistance for all cases below the  $AUC_{0-24}/MIC$  ratio breakpoint was 7 days versus 6 days for those cases above the breakpoint. This difference was not significant, irrespective of treatment or organism. Figure 2 is a Kaplan-Meier plot of the probability of remaining susceptible over time, from the initiation of therapy. The three curves represent three distinct groups: (i) the cases fit by the model below the  $AUC_{0-24}/MIC$  breakpoint of 100 ( $n = 17$ ); (ii) the cases consistent with type I  $\beta$ -lactamase-producing organisms treated with  $\beta$ -lactam monotherapy, which did not exhibit an  $AUC_{0-24}/MIC$  relationship ( $n = 14$ ); and (iii) the cases fit by the model above the  $AUC_{0-24}/MIC$  breakpoint of 100 ( $n = 97$ ). For the organisms represented by curve i ( $n = 17$ ), the times to selection of 25, 50, and 75% resistance occurred by days 5, 10, and 14, respectively. For the

gram-negative rods represented by curve ii ( $n = 14$ ), the time to selection of 25% resistance occurred by day 5, and the time to selection of 50% or greater resistance occurred by day 16. For all other organisms, the cumulative rate of resistance development was approximately 9% and remained relatively consistent over time, with all cases of resistance occurring by day 13. A statistically significant difference was noted between groups 1 and 3 ( $P < 0.001$ ) and between groups 1 and 2 ( $P < 0.001$ ). Groups 2 and 3 did not differ ( $P = 0.322$ ). However, the numbers of cases in these two groups are small and a type II error may exist. For these two groups, which exhibit different relationships of antimicrobial exposure to response, the rates of selection of bacterial resistance are similar.

## DISCUSSION

In this population of acutely ill patients with LRTIs, there was an inverse-effect relationship between the probability of

TABLE 5. Univariate analysis results of resistance development by patient risk factors and characteristics

Risk factor or characteristic <sup>a</sup>	P <sup>b</sup>
Age .....	0.16 <sup>c</sup>
Weight.....	0.17 <sup>c</sup>
$AUC_{0-24}/MIC$ ratio .....	$\ll 0.001^c$
Gender .....	0.77
Infection with:	
<i>Pseudomonas</i> spp. ....	0.0002
Type I $\beta$ -lac <sup>+</sup> organisms.....	0.74
Other gram-negative rods.....	0.008
Miscellaneous organisms .....	0.04
Treatment with:	
CIP + PIP .....	0.11
CAZ.....	0.76
Cefmenoxime .....	0.01
CIP .....	0.54
CAZ + TOB.....	0.29
Ventilator dependent .....	0.91
Major surgery.....	0.24
COPD.....	0.67
Steroids .....	0.82
Diabetes mellitus .....	0.82
Malignancy .....	0.22
Chemotherapy or radiation therapy.....	0.57
Prior antibiotics.....	0.02
Type I $\beta$ -lac <sup>+</sup> organisms; $\beta$ -lactam monotherapy .....	0.001

<sup>a</sup>  $\beta$ -lac<sup>+</sup>,  $\beta$ -lactamase producing; CIP, ciprofloxacin; PIP, piperacillin; CAZ, ceftazidime; TOB, tobramycin.

<sup>b</sup> As evaluated by the chi-square test.

<sup>c</sup> Determined by Kruskal-Wallis one-way analysis of variance.



TABLE 6. Summary statistics for  $AUC_{0-24}/MIC$  ratios by organism groups and by treatment groups

Organism or treatment group <sup>a</sup>	$AUC_{0-24}/MIC$		
	Median	Mean (SD)	Range
<b>Organism group</b>			
1			
Susceptible	274	735 (1,501)	73–6,967
Resistant	74	145 (190)	15–733
2			
Susceptible	667	1,604 (1,720)	33–5,541
Resistant	693	3,517 (5,042)	217–14,190
3			
Susceptible	2,396	4,849 (8,510)	169–45,263
Resistant	290	379 (399)	48–887
4			
Susceptible	843	5,937 (15,970)	130–56,423
<b>Therapy group</b>			
<b>CIP/PIP</b>			
Susceptible	302	1,130 (1,693)	118–6,103
<b>CAZ</b>			
Susceptible	5,303	13,346 (19,174)	218–56,423
Resistant	887	5,246 (7,746)	661–14,190
<b>CEFMNX</b>			
Susceptible	668	1,999 (2,305)	73–6,852
Resistant	278	1,655 (3,259)	18–9,822
<b>CIP</b>			
Susceptible	582	1,516 (1,904)	33–8,158
Resistant	72	185 (275)	15–1,024
<b>CAZ/TOB</b>			
Susceptible	2,499	3,069 (3,075)	199–10,950
Resistant (n = 1)	733		

<sup>a</sup> Organism groups are as defined in Materials and Methods. CIP/PIP, ciprofloxacin plus piperacillin; CAZ, ceftazidime; CEFMNX, cefmenoxime; CIP, ciprofloxacin; CAZ/TOB, ceftazidime plus tobramycin.

the development of bacterial resistance and the  $AUC_{0-24}/MIC$  ratio. This relationship was strongest for *Pseudomonas aeruginosa* treated with ciprofloxacin, but was also found within other organism groups and antibiotic treatments. These findings support previous reports of the selection of resistance within *Pseudomonas aeruginosa* strains from studies utilizing in vitro pharmacodynamic models with various ciprofloxacin dosing regimens (18, 19). In the in vitro model studies, bacterial re-

TABLE 7. Pharmacodynamic model goodness of fit

Antimicrobial exposure ( $AUC_{0-24}/MIC$ ) <sup>a</sup>	n	% Observed resistance	% Modelled fits
<100	17	82.4	82.6
≥100	97	9.3	9.2
0–50	8	75.0	82.6
50–100	9	88.9	80.9
100–250	21	14.3	10.2
250–500	16	6.3	9.2
500–1,000	18	22.2	9.2
>1,000	42	2.4	9.2
BLAC, all	14	64.3	64.3
BLAC, 100–250	1	100	83.2
BLAC, 250–500	3	66.7	76.6
BLAC, 500–1,000	4	75.0	72.3
BLAC, >1,000	6	50.0	49.6

<sup>a</sup> BLAC, type I  $\beta$ -lactamase-producing organisms treated with  $\beta$ -lactam monotherapy. The numbers following BLAC indicate the  $AUC_{0-24}/MIC$  ranges; all, all  $AUC_{0-24}/MIC$  values.

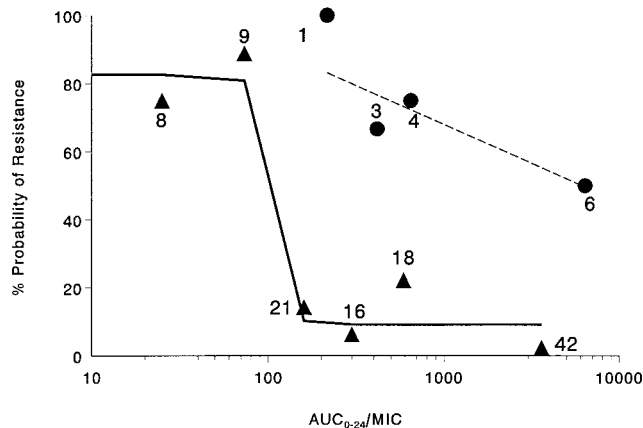


FIG. 1. Relationship between the percent probability of resistance and the antimicrobial exposure as expressed by the average daily  $AUC_{0-24}/MIC$  ratio. ●, observed data for type I  $\beta$ -lactamase-producing gram-negative rods treated with  $\beta$ -lactam monotherapy; ▲, observed data for all other cases. The solid line represents the modelled response surface for all cases, classified as either susceptible or resistant, within the data set for which the  $AUC_{0-24}/MIC$  inverse-effect relationship applied. The dashed line represents the line of best fit for the group 2 organisms treated with  $\beta$ -lactam monotherapy. Observed cases identified by symbols and numbers are plotted as single points within an  $AUC_{0-24}/MIC$  ratio category at the median value for that category.

sistance did not occur when a dose of 1,200 mg of ciprofloxacin was administered once daily, but it was associated with a  $C_{max}/MIC$  ( $C_{max}$ , maximum concentration of the drug in serum) ratio of <10 (19). In a similar in vitro model, the investigators were able to select subvariant bacterial populations with increasing MICs related to a  $C_{max}/MIC_{50}$  ( $MIC_{50}$ , MIC at which 50% of isolates are inhibited) ratio of 7.3 and a breakpoint  $AUC/MIC_{50}$  ratio of 95  $SIT^{-1} h$  (18), essentially identical to our  $AUC_{0-24}/MIC$  breakpoint of 100. These results suggest that the  $AUC_{0-24}/MIC$  ratio may be a useful parameter to guide therapy with the goal of preventing the selection of resistance (18).

In a neutropenic rat model of *Pseudomonas* sepsis, Drusano and colleagues studied lomefloxacin and found that either the  $C_{max}/MIC$  ratio or the  $AUC/MIC$  ratio was a significant predictor of survivorship (11). However, when the  $AUC$  was held constant but the dose was changed to produce very high  $C_{max}/$

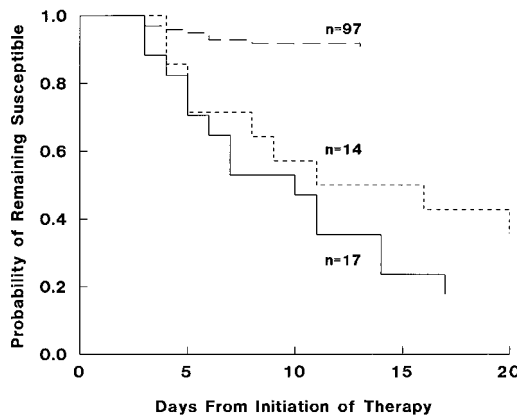


FIG. 2. Relationship between the probability of the development of resistance and treatment duration (days). Solid line, data for cases in which the  $AUC_{0-24}/MIC$  ratio was <100; small dashed line, data for type I  $\beta$ -lactamase-producing gram-negative rods treated with  $\beta$ -lactam monotherapy; large dashed line, data for cases in which the  $AUC_{0-24}/MIC$  ratio was  $\geq 100$ .

MIC ratios ( $\geq 20:1$ ), outcomes were significantly improved. Success in this model may reflect successful eradication of bacterial subpopulations that remain viable and that are selected for resistance by lower exposures. Unfortunately, dosing regimens of the fluoroquinolones which could be tolerated in human patients would not achieve these very high  $C_{\max}/\text{MIC}$  ratios for *Pseudomonas*. Finally, our data provide support for the observation that a  $C_{\max}/\text{MIC}$  ratio of approximately 5:1 may correspond to an  $\text{AUC}_{0-24}/\text{MIC}$  ratio of 100 for ciprofloxacin.

We agree with Madaras-Kelly (18), that the  $\text{AUC}_{0-24}/\text{MIC}$  ratio is the most reasonable parameter, as it combines both concentration intensity and exposure over time. In addition, we do not advocate  $C_{\max}/\text{MIC}$  targets over  $\text{AUC}/\text{MIC}$  targets because the dosing regimens suggested by Marchbanks and Drusano to achieve their effective  $C_{\max}/\text{MIC}$  ratios are not achievable in human pseudomonas infections, without unacceptable side effects. The other problem is that unlike  $\text{AUC}_{0-24}/\text{MIC}$  ratios,  $C_{\max}/\text{MIC}$  ratios are not additive. Thus, it is not possible to optimize two antibiotics based on the  $C_{\max}/\text{MIC}$  ratio.

A recent review investigating the frequency of bacterial resistance included 173 clinical trials, incorporating 14,000 patients and seven antibiotic classes (13). One study included within our data set was among the 173 trials reviewed (22).

The authors reported an overall rate of bacterial resistance development of 4%, and in LRTI patients the incidence was 8.9%. The rates of resistance development reported for the intensive-care unit (ICU) population (7%) and the mechanically ventilated population (9%) were less than the 25% overall rate of resistance development reported in our study. However, our patients were primarily ICU patients and 72% were mechanically ventilated. Factors within the ICU setting such as mechanical ventilation and multiple underlying diseases appeared to contribute to bacterial resistance. Our observation of a lower frequency of resistance with combination therapy was also consistent with the finding of the review. Combination regimens had median  $\text{AUC}_{0-24}/\text{MIC}$  ratios similar to those of the monotherapy regimens (Table 6), but the minimum values more frequently exceeded the  $\text{AUC}_{0-24}/\text{MIC}$  ratio of 100. When two antibiotics are used, there is always a greater chance that one of the two may have an  $\text{AUC}_{0-24}/\text{MIC}$  ratio above 100, thus explaining the better overall protection from resistance of combination therapy in pharmacodynamic terms.

The organisms which failed to demonstrate a statistically significant relationship between the development of resistance and the  $\text{AUC}_{0-24}/\text{MIC}$  ratio were all  $\beta$ -lactamase-producing organisms treated with  $\beta$ -lactam monotherapy. Type I  $\beta$ -lactamase production is readily selected by the broad-spectrum cephalosporins in *Enterobacter* spp. and is associated with the development of resistance by the simple eradication of the susceptible subpopulations (3, 6, 20). The selection of resistant mutants during therapy results in a greater proportion of organisms producing adequate quantities of enzyme, which readily hydrolyzes most  $\beta$ -lactams (20), and the net effect is a rise in MIC. We suggest that in this situation there are bacterial subpopulations with baseline enzyme production present in the culture which will not be eradicated even with exposure of the entire population above the  $\text{AUC}_{0-24}/\text{MIC}$  ratio of 100. Although the initial MICs seem to be low, they quickly increase as the susceptible organisms are eradicated leaving the resistant mutants behind. These findings provide further argument to avoid the use of monotherapy with broad-spectrum cephalosporins in these organisms, since our data show that variable rates of selection of bacterial subpopulations cannot be predicted by baseline MIC testing. Combination therapy with an antibiotic not susceptible to this mechanism is the necessary

procedure. We surmise that an  $\text{AUC}_{0-24}/\text{MIC}$  ratio above 100 is required for the second (non- $\beta$ -lactam) drug, or the probability for failure will also be increased for the combination regimen. This may explain why aminoglycosides added to broad-spectrum cephalosporins did not always prevent the selection of resistance in *E. cloacae* (3), since aminoglycosides alone do not often produce  $\text{AUC}_{0-24}/\text{MIC}$  ratios above 100.

A mechanistic discussion of how  $\text{AUC}_{0-24}/\text{MIC}$  relationships might explain the selection of each type of resistance during therapy is clearly beyond the scope of this paper. However, the mechanism of resistance in many of our patients is most likely the selection of preexisting subpopulations of resistant mutants. In gram-negative pathogens the mutant subpopulations may have alterations in porin channels, plasmids containing resistant genes, active efflux pumps, or mutations in regulatory genes controlling  $\beta$ -lactamase production (6, 15, 20). In each of these cases, the action of the antibiotic is to eliminate the susceptible majority, leaving the selected remainder intact. In the absence of host defense or in the presence of foreign surfaces, the selected remainder quickly becomes the dominant population.

This study has several limitations. First, the retrospective nature of the review and the potential for bias in selection of resistant cases must be acknowledged, as ICU patients with nosocomial pneumonia are frequently colonized with gram-negative organisms and many have received previous courses of antimicrobial therapy. These data are not epidemiologic study data useful for determining the comparative frequency of resistance across patient populations, among ICUs, or among different antibiotics. Second, controversy will continue over the relative value of the different methods of lower respiratory tract specimen sampling and culture. The patients were clinically similar in all four trials (Table 1), and all the patients were treated in one institution under observation by the same investigators. There were common inclusion criteria for all of our nosocomial pneumonia trials, and most cases reported here were eventually used as part of new drug application submissions. Third, this study assumed that the resistant organisms isolated during therapy and follow-up were the same organisms as those isolated in the original cultures. Molecular typing (DNA) methods would be required to compare strain homologies, and these tests were not performed. Thus, the impact of  $\text{AUC}_{0-24}/\text{MIC}$  ratios below 100 on selection versus the impact of new mutations or the introduction of new organisms cannot be definitively resolved. We can state with statistical certainty that when the  $\text{AUC}_{0-24}/\text{MIC}$  ratio is below 100, 82.4% of the organisms developed resistance via some mechanism, the most likely being the selection of a subpopulation. When the  $\text{AUC}_{0-24}/\text{MIC}$  ratio was above 100, only 9% of similar patients with similar organisms developed resistance (Table 7 and Fig. 2).

These results provide clinical data to support previously reported findings from in vitro models and from animal studies. They demonstrate a relationship between the antimicrobial pharmacodynamic parameter  $\text{AUC}_{0-24}/\text{MIC}$  and the development of bacterial resistance. The association between resistance and bacterial MIC "coverage" by drug concentration is clearly stronger than any of the previously elucidated clinical risk factors. This finding is intuitively logical, since exactly the same conclusion arises from studies conducted in vitro and with animal models in the absence of patient factors and since the nosocomial pneumonia patient represents one of the situations where there is relatively little contribution of host defense to outcome.

Our results also suggest resistance can also be avoided with attention to dosing, since dosing regimens which provide an

$AUC_{0-24}/MIC$  ratio of at least 100 appear to reduce the rate of the development of bacterial resistance in acutely ill patients with nosocomial pneumonia. Additional molecular studies of subpopulations will be needed to identify the relationship between  $\beta$ -lactam monotherapy and the selection of resistant organisms consistent with type I  $\beta$ -lactamase producers. Finally, clinical trials evaluating new antimicrobial agents should measure antimicrobial exposure parameters, such as the  $AUC_{0-24}/MIC$  ratio, in order to relate these findings to both microbiologic and clinical outcomes and to determine the relationship between these indices and the development of antibacterial resistance.

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