Susceptibility Testing Accuracy of an Enterobacteriaceae Population Enriched for CTX-M-type ESBL -producing Isolates (China): Intermethod Analysis for Nine Important Parenteral B-lactams

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RESULTS

ABSTRACT

Background: To assess the wide geographical applicability of the current (C) or proposed (P) susceptibility (S) breakpoint (BP) criteria for 9 ß-lactams, a collection of Enterobacteriaceae (ENT) enriched with CTX-M-type ESBL-producing isolates from China recent clinical isolates (blood cultures. 68.6%) of ENT were analyzed: E. coli (100), Klebsiella spp. (60), Proteus spp.

. (43). Serratia spp. (31) and others (68). One-in-five (21.5%) produced an ESBL with predominance of CTX M-type enzymes (94.7%: endemic types 3. 14. 24 and 28). Isolates were tested by CLSI broth microdilution and disk diffusion (DD) methods. Categorical agreement (CA) using MIC C-BP and those based on PK/PD target attainment (90% at MIC, T > 50%) & Monte-Carlo simulations (P-BP) were analyzed. The intermediate category was removed for cefoxitin and cefuroxime with no change in S-BP because of MIC distributions of wild-type S populations **Results:** The error rates and CA results are summarized in the Table.

	S P-BP	False-S	False-R	CA
Antimicrobial agent	MIC (µg/ml)/disk (mm)	C-BP/P-BP	C-BP/P-BP	C-BP/P-BP
Aztreonam	≤4/≥21	0.0/0.0	0.0/0.0	93.2/97.5
Cefepime	≤4/≥21	0.0/0.3	0.0/0.0	95.2/96.3
Cefotaxime (CT)	≤1/≥26	0.0/0.0	0.3/0.0	92.6/99.2
Cefoxitin	<u><</u> 8/≥18	0.8/7.3	0.0/1.1	85.1/91.6
Ceftazidime	≤4/≥21	0.0/0.0	0.0/0.0	97.2/98.0
Ceftriaxone (TX)	≤1/≥23	0.0/0.0	0.3/0.0	94.6/99.2
Cefuroxime (XM)	≤8/≥18	0.0/1.7	0.0/1.1	96.0/97.2

Regression analysis showed (r) values ranging from 0.84 (cefotetan [CN]) to 0.98 (CT, XM, TX). The MIC P-BP of \leq 2 µg/ml for CN and ceftizoxime with corresponding DD correlates, \geq 24 and \geq 23 mm, resulted in only 7.6 and 5.1% error rates. Conclusions: Our results confirm that C and P MIC BP criteria with appropriately selected DD correlates have acceptable, generall improved CA when tested against ENT enriched for CTX-M-type ESBL-producing strains endemic in numerous locations outside

INTRODUCTION

Extended-spectrum ß-lactamases (ESBLs) are potent plasmid-mediated enzymes which can hydrolyze oxyimino-cephalosporins and monobactams. These Bush group 2be ESBL enzymes, among others, constitute a heterogeneous molecular cluster comprised of βlactamases sharing 20 to > 99% identity. Based on their substrate preferences, these enzymes are broadly classified as ceftazidimases (preferential hydrolysis of ceftazidime > cefotaxime) and cefotaximases (preferential hydrolysis of cefotaxime > ceftazidime). The well known TEM and SHV ESBL groups (> 130 members) described worldwide are predominantly ceftazidimases. Recently, non-TEM and non-SHV plasmid-mediated ESBL types have been reported both in the ceftazidimases and cefotaximases groups. The cefotaximases include SFO-1, BES-1 and CTX-M enzymes. The diversity of molecular structure and hydrolytic spectra of the ESBL enzymes, in addition to the differences of the geographical occurrence of the individual ESBL-types contribute to the challenges in designing appropriate laboratory tests to detect the presence of these enzymes or to guide accurate therapeutic decisions and infection control interventions

The current Clinical and Laboratory Standards Institute (formerly the National Committee for Clinical Laboratory Standards [NCCLS]) recommendation for ESBL screening and confirmatory tests of *E. coli* and *Klebsiella* spp. was established in response to the reports of therapeutic failures due to ESBL-producing strains treated with cephalosporins which were tested as susceptible by CLSI methods. However, these recommendations do not address the entire issue for detection of the presence of ESBLs in all species of the Enterobacteriaceae and also require qualifying reports to the physicians. To ensure recommendations that predict favorable clinical response when testing contemporary isolates of Enterobacteriaceae, the standardizing authorities (NCCLS and others) have recognized the need to re-evaluate the existing breakpoints and consider alternative susceptible criteria for all clinically relevant cephalosporins

Studies using Monte-Carlo simulations and pharmacokinetic/pharmacodynamic (PK/PD) target attainment (≥ 90.0%) data support the adjustment of susceptibility MIC breakpoints that are two- to 16-fold lower for aztreonam, cefepime, cefotaxime, cefotetan, ceftazidime, ceftriaxone and ceftizoxime. For cefuroxime and cefoxitin based on the MIC population distributions and target attainment results, removal of the intermediate category accompanied by no change in the susceptible breakpoints has been proposed. Earlier reports have clearly demonstrated that both the current and the proposed alternative MIC breakpoints and corresponding disk diffusion criteria have an acceptable intermethod error rate when testing an ESBL-enriched collection of Enterobacteriaceae from the Americas against cephalosporin and monobactam antimicrobials. However, ESBLs differ in their substrate specificity and there are well recognized differences in the geographical occurrence of the various ESBL-types. To assess the global applicability of these breakpoint criteria, particularly outside of the United States (USA), we have reassessed the intermethod error rates of current and the proposed breakpoint criteria for a collection of Enterobacteriaceae selected for a predominantly CTX-M-type ESBL-producing population, endemic in China and the Far East, but becoming worldwide in range.

MATERIALS AND METHODS

Strain collection. A collection of 354 strains of Enterobacteriaceae isolated from clinical specimens in 2003 - 2004 were analyzed. The collection was comprised of the following species (strains): Escherichia coli (100), Klebsiella spp. (60), Enterobacter spp. (43), Serratia spp.(31), Citrobacter spp. (29), Proteus mirabilis (28), indole-positive Proteae (24), Shigella spp.(15), Salmonella spp. (10) and others (14). Specimen type data was available for 83.9% (297) of the isolates. The vast majority of strains were from bloodstream infections (243; 81.8%), with the remaining organisms isolated from urine (24; 8.1%), gastrointestinal tract (22; 7.4%), documented skin and soft tissue infections (five; 1.7%) and respiratory specimens from patients with nosocomial pneumonia (three; 1.0%). More than one in five isolates (76; 21.5%) were confirmed ESBL-producing strains comprised of E. coli (48), K. pneumoniae (22), K. oxytoca (three), and one strain each of *P. mirabilis*, *P. vulgaris* and *Providencia stuartii*. The type of ESBL produced was identified by PCR amplification and gene sequencing procedures for 75.0% (57 of 76) of strains. These isolates were obtained from China and the CTX-M group was the dominant (94.7%; 54 of 57) ESBL-type; containing CTX-M-14 (29), CTX-M-3 (10), CTX-M-24 (five), CTX-M-3 + CTX-M-14 (five), CTX-M-28 (three), CTX-M-14 + SHV-12 (two), with SHV-type ESBLs constituting the remainder of isolates (SHV-12 [two] and SHV-5 [one]).

Susceptibility testing methods. Minimal inhibitory concentrations (MICs) of nine B-lactams were determined by reference broth microdilution method, and interpreted according to CLSI criteria. MIC results were also determined using the Etest strips in accordance with the manufacturer's recommendations (AB BIODISK, Solna, Sweden). Simultaneous disk diffusion testing was performed for all B-lactams analyzed and interpreted in accordance with the CLSI guidelines. The B-lactams analyzed included: aztreonam, cefepime, cefotaxime, cefotetan, cefoxitin, ceftazidime, ceftriaxone, ceftizoxime and cefuroxime.

Selection of proposed MIC breakpoints: Using published mean PK parameter estimates and dispersion measures from volunteer studies, and PK-PD targets derived from a murine infection model (% free-drug time > MIC₅₀), Monte-Carlo simulation (10,000 subjects) was used to evaluate the probability of PK/PD target attainment (≥ 90.0%) for standard β-lactam dosing regimens. Simulation results supported the reduction of some susceptibility MIC breakpoints two- to eight-fold for aztreonam, cefepime, cefotaxime, cefotetan ceftazidime, ceftizoxime and ceftriaxone. Current susceptibility breakpoints were noted to bisect the wild-type MIC population distribution for cefoxitin. It was also evident that susceptibility breakpoints bisecting MIC population distributions could be avoided assuming the use of higher labeled doses for serious infections for cefuroxime. For both cefuroxime and cefoxitin, proposed criteria include removal of the intermediate category with no modification in the susceptible breakpoint. Correlate disk diffusion criteria were recommended corresponding to the proposed changes in the MIC breakpoints. Current MIC and disk diffusion breakpoint criteria were applied as published by the CLSI document M100-S15.

Analysis of the results: Results of the disk diffusion tests were compared with those of the reference broth microdilution supported by the Etest (AB BIODISK, Solna, Sweden) method. The data was plotted as scattergrams and the least squares method was used to calculate the regression equation. Intermethod-error rates were calculated, and the results were compared to an earlier experiment for cefepime, used as an internal control. The criteria used to calculate the inter-method agreement for MIC and disk diffusion testing included the current CLSI criteria for the nine B-lactams, and the proposed (generally lower) interpretive criteria.

Table 1. Susceptibility profiles of nine ß-lactams tested against 354 Enterobacteriaceae by reference/standardized NCCLS methods. MIC_{50/90} (% susceptible)^b Ceftriaxone Cefuroxime Ceftizoxime Aztreonam 32/>256(45) 0.25/>256(52) 0.12/16(80) 0.12/>256(57) 8/>256(53) Klebsiella spp. (60) Citrobacter spp. (29 P. mirabilis (28 128/>256(33) 0.03/32(88) Indole-positive Proteae (2)

• P. mirabilis, Shigella spp. and Salmonella spp. had the highest susceptibility rates (ranging from 90.0 to 100.0%) compared to the other species of Enterobacteriaceae when tested against the ß-lactams (Table 1). For E. coli and Klebsiella spp. containing the majority of ESBL- producing strains (96.1%; 73 of 76 strains), the susceptibility rates for the ESBL screening compounds (cefotaxime, ceftriaxone, ceftazidime and aztreonam) ranged from only 52.0 to 90.0%. Noteworthy for these two species, was the lower susceptibility rates to ceftriaxone (\leq 57.0%) and cefotaxime (\leq 72.0%) compared to the susceptibility rates to ceftazidime (≥ 87.0%); characteristics of CTX-M-type enzymes.

a. Only the species or genus groups with \geq 10 strains were tabulated

b. Susceptibility as defined by CLSI excluding ESBL screening criteria.

- The scattergrams for the entire collection of 354 Enterobacteriaceae including the 76 ESBL-producing strains tested against the nine ß-lactams are plotted in Figures 1 through 9. Regression statistics showed excellent correlation between the two methods for most of the antimicrobials analyzed, with the regression coefficient ranging from r = 0.84 (cefotetan) to r = 0.98 (cefotaxime, cefuroxime and ceftriaxone).
- The intermethod categorical concordance for the ß-lactams analyzed remained acceptable for the CTX-M-type ESBL enriched Enterobacteriaceae isolates analyzed in the present study when applying the CLSI interpretive criteria for eight of the nine B-lactams analyzed (except cefoxitin), with an absolute intermethod agreement ranging from 92.6 (cefotaxime) to 97.8% (ceftazidime), see Table 2 and Figures 1-9.
- Very major errors (false-susceptible) were nil (0.0%) for seven of the nine B-lactams. Major errors (falseresistant) were absent for all B-lactams, except cefotaxime, cefotetan and ceftriaxone (0.3%). Minor error rates ranged from 0.8% for cefotetan to the highest rate of 14.1% noted for cefoxitin, resulting in a low intermethod concordance rate of 85.1% overall.

ble 2.	Intermethod error rates for 354 Enterobacteriaceae isolates based on current CLSI interpretive
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	Method	Current criteria ^a		Error rates ^b				
Antimicrobial agent		S	I	R	Very major	Major	Minor	Intermethod agreement
Aztreonam	MIC Disk	≤8 ≥22	16 16-21	≥32 ≤15	0.0	0.0	6.8	93.2
Cefepime	MIC Disk	≤8 ≥18	16 15-17	≥32 ≤14	0.0	0.0	4.8	95.2
Cefotaxime	MIC Disk	≤8 ≥23	16-32 15-22	≥64 ≤14	0.0	0.3	7.1	92.6
Cefotetan	MIC Disk	≤16 ≥16	32 13-15	≥64 ≤12	1.1	0.3	8.0	97.8
Cefoxitin	MIC Disk	≤8 ≥18	16 15-17	≥32 ≤14	0.8	0.0	14.1	85.1
Ceftazidime	MIC Disk	≤8 ≥18	16 15-17	≥32 ≤14	0.0	0.0	2.8	97.2
Ceftriaxone	MIC Disk	≤8 ≥21	16-32 14-20	≥64 ≤13	0.0	0.3	5.1	94.6
Ceftizoxime	MIC Disk	≤8 ≥20	16 15-19	≥32 ≤14	0.0	0.0	3.4	96.6
Cefuroxime	MIC Disk	≤8 ≥18	16 15-17	≥32 ≤14	0.0	0.0	4.0	96.0

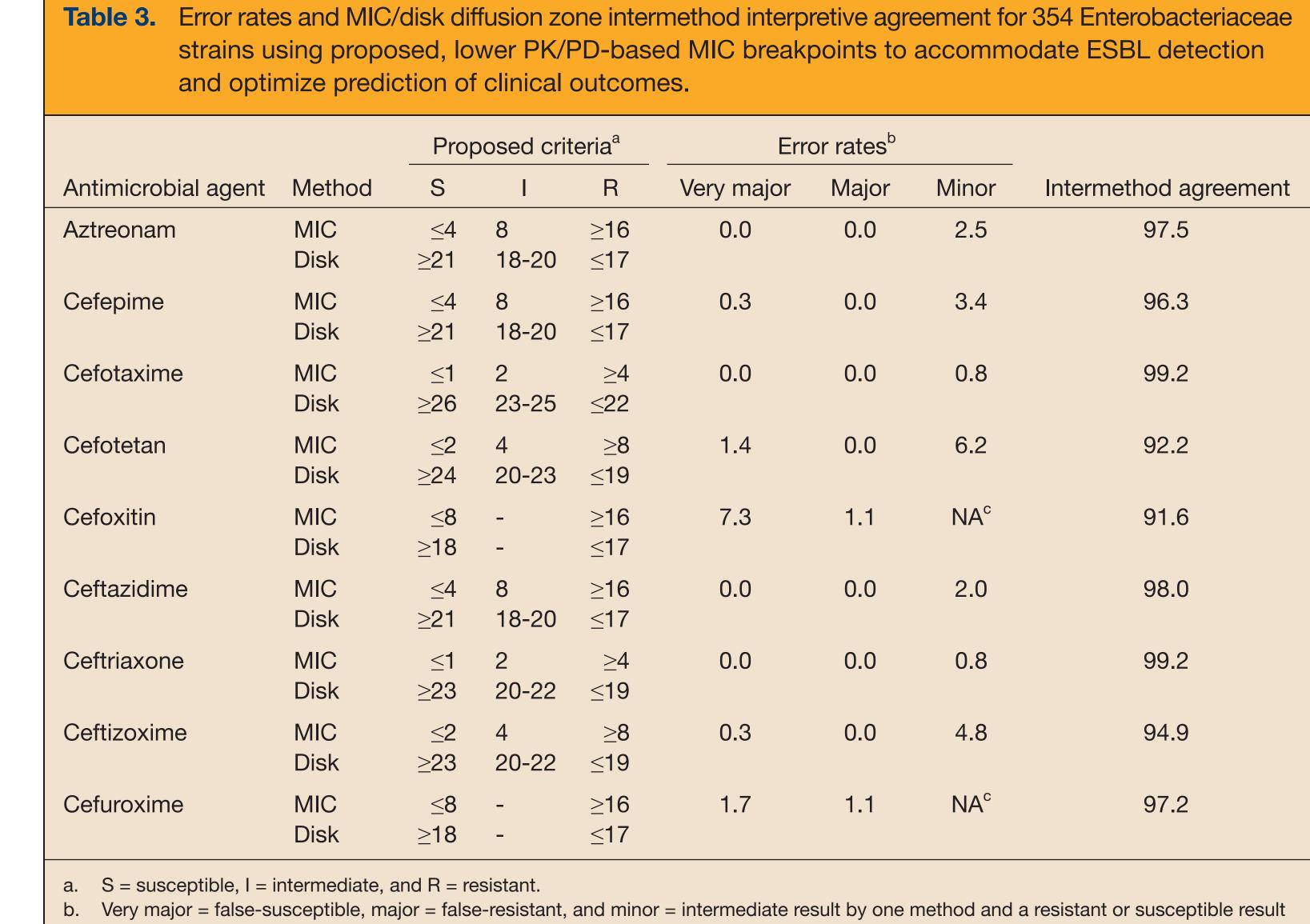
o. Very major = false-susceptible, major = false-resistant, and minor = intermediate result by one method and a resistant or susceptible result

with the other test.

The intermethod concordance rates for the modified breakpoint criteria analyzed using an Enterobacteriaceae collection enriched with CTX-M-type ESBL-producing strains (Table 3 and Figures 1-9) was very acceptable, with intermethod agreement rates ranging from 91.6 (cefoxitin) to 99.2% (ceftriaxone and cefotaxime). The intermethod agreement improved for seven of the nine \(\textit{B-lactams} \) analyzed; aztreonam (+ 4.3%), cefepime (+ 1.1%), cefotaxime (+ 6.6%), cefoxitin (+ 6.5%), ceftazidime (+ 0.8%), ceftriaxone (+ 4.6%) and cefuroxime (+1.2%).

- While the proposed interpretive criteria for cefoxitin and cefuroxime (no change in the susceptible breakpoints and elimination of the intermediate category) resulted in an increase in significant errors (verymajor [+ 1.7% cefuroxime; + 6.5% cefoxitin] and major [+ 1.1%] error) for both antimicrobials) the intermethod categorical agreement remained at acceptable levels of 91.6% (cefoxitin) and 97.2%
- Noteworthy was the absence of both very major (false-susceptible) and major (false-resistant) errors for

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 The proposed (generally lower) MIC breakpoints were selected using a rigorous scientific process, taking into consideration the PK/PD data and Monte-Carlo simulations, as well as the use of MIC population analyses. These modified breakpoints will assist clinicians in the choice of an appropriate antimicrobial agent based on the in-vitro activity

antimicrobials analyzed.

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putative resistance mechanism to unaffected peer drugs.

against the pathogen in question, and not by extrapolation of the

CONCLUSIONS

disseminated in the last decade to constitute the most widespread

potent enzymes compared to the other ESBL types include: 1.)

group of ESBL enzymes worldwide. The distinctive features of these

emergence not limited to nosocomial strains of E. coli and Klebsiella

spp. with detection in community bacterial strains of Vibrio cholerae,

non-typhoidal Salmonella spp. and Shigella spp.; 2.) identification of

potential reservoirs of the resistant determinants in commensal *E. coli*

emergence linked to the widespread use of cefotaxime and ceftriaxone.

CLSI interpretive criteria, the intermethod (MIC versus zone diameter)

error rates continue to be acceptable (92.6 to 97.8%) for eight of the

nine antimicrobials analyzed. The new susceptible breakpoints also

99.2%), with significant improvements (+ 1.1% for cefepime to + 6.6%

had acceptable intermethod error rates for all B-lactams (91.6 to

for cefotaxime) in the categorical agreement for nearly of the

isolates from healthy children and animals; 3.) rapid dissemination

resulting in small or countrywide or international outbreaks; and 4.)

Our results with CTX-M type strains indicates that using the current

CTX-M type enzymes are cefotaximases which have rapidly

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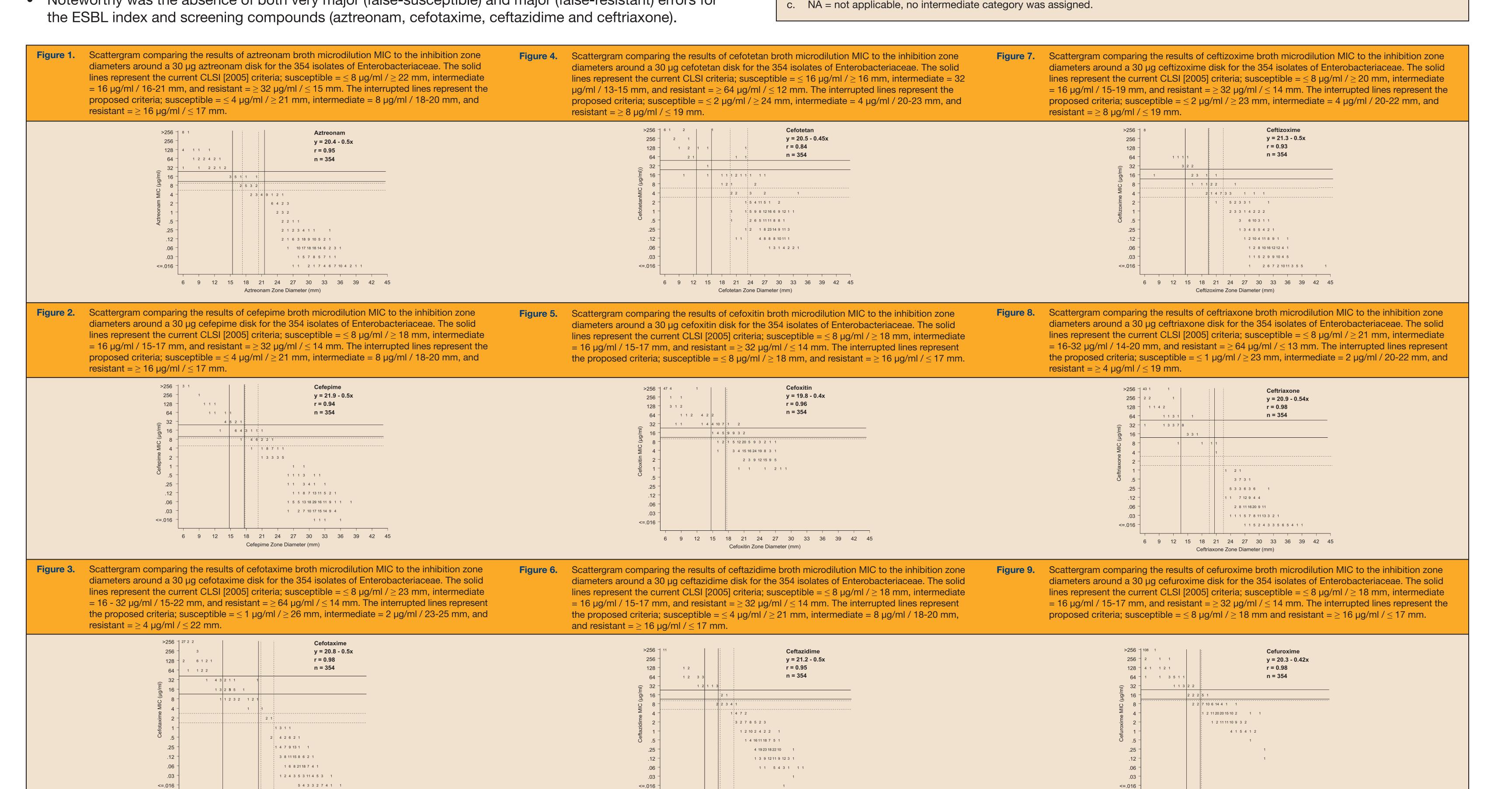
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