Evaluation of the Sirscan Automated Zone Reader in a Clinical Microbiology Laboratory

ANTONE A. MEDEIROS* AND JOYCE CRELLIN

Brown University, Lifespan Academic Medical Center, The Miriam Hospital, Providence, Rhode Island 02906

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We compared readings of Kirby-Bauer plates by the Sirscan, an automated image analyzer that measures zone diameters, to those of experienced clinical microbiologists measuring zones with a hand-held caliper interfaced to a computer and with a ruler. To read plates of Escherichia coli, Morganella morganii, and Pseudomonas aeruginosa containing 12 antibiotic disks the Sirscan took 11 s; technologists took 28 s by caliper and 39 s by ruler. Reading times of four different technologists ranged from 22 to 44 s with the caliper and 10 to 12 s with Sirscan. Upon repeated testing zone size variation rarely exceeded 3 mm by caliper and 1 mm by Sirscan. Over a 4-month period, 368 clinical isolates were tested prospectively by both methods in the Clinical Microbiology Laboratory of the Miriam Hospital. There was good correlation of zone sizes for most antibiotics, but Sirscan zone diameter measurements tended to be 3 to 5 mm larger than caliper readings for ciprofloxacin, norfloxacin, aztreonam, erythromycin, clindamycin, and trimethoprim-sulfamethoxazole. Very major errors (resistant by caliper and susceptible by Sirscan) occurred with 10 of 3,770 readings (0.3%), mainly where breakpoint criteria lacked an intermediate zone. They occurred in testing staphylococci with amoxicillinclavulanate (5 of 127 isolates, 3.9%), pseudomonas with piperacillin (1 of 28, 3.6%), coagulase-negative staphylococci with oxacillin (2 of 74, 2.7%), gram-negative bacilli with cefuroxime (1 of 209, 0.5%), and mixed species with trimethoprim-sulfamethoxazole (1 of 366, 0.3%). The Sirscan zone reader facilitates accurate, fully quantitative susceptibility testing in clinical microbiology laboratories.

Early detection of emerging resistance mechanisms requires quantitative susceptibility testing, either zone diameter measurements or full panel MICs, as recommended by the ASM Task Force on Antibiotic Resistance (2). However, most commonly used automated methods provide only breakpoint values of antibiotic susceptibility. Single disk diffusion provides more endpoint values, about 35, than most full-panel dilution methods, about 12, but measuring zone sizes is tedious, timeconsuming, and fraught with transcription errors. An automated method of measuring zone sizes would obviate these limitations.

In our clinical microbiology laboratory, we evaluated the Sirscan (i2a, Montpelier, France), an automated image analyzer that measures zone diameters and provides a user-programmed expert system that screens the results of each isolate. The program also extrapolates zone sizes to MICs and reports both.

Susceptibility tests. Testing was performed by the Kirby-Bauer single disk diffusion method according to National Committee on Clinical Laboratory Standards (NCCLS) guidelines (4), using 150-mm round plates of Mueller-Hinton agar purchased from BBL, Becton-Dickinson, Cockeysville, Md.

Strains tested. Strains were fresh clinical isolates from the Clinical Microbiology Laboratory of the Miriam Hospital. The number of isolates tested per species are as follows: 101 isolates of *Escherichia coli*, 74 isolates of coagulase-negative staphylococci, 53 isolates of *Staphylococcus aureus*, 28 isolates of *Pseudomonas aeruginosa*, 25 isolates of *Klebsiella pneumoniae*, 15 isolates of *Serratia marcescens*, 13 isolates of *Kleb*

siella oxytoca, 13 isolates of Proteus mirabilis, 12 isolates of Enterobacter cloacae, 8 isolates of Citrobacter freundii, 8 isolates of Acinetobacter baumanii, 8 isolates of S. maltophilia, 7 isolates of Enterobacter aerogenes, and 1 isolate each of Serratia liquefaciens, Proteus penneri, and Salmonella sp. Thirty-five percent were urine isolates, 21% were from sputum, 19% were from blood, 11% were from wounds, and the remainder came from miscellaneous sources.

Control strains were *E. coli* ATCC 25922, *P. aeruginosa* ATCC 27853, and *S. aureus* ATCC 25923.

Zone diameter readings with a ruler. A technologist experienced in reading zones with a ruler placed the Kirby-Bauer (K-B) plate (150-mm diameter and round) on a view box with reflected light against a black background and measured the zone diameters manually. She dictated the readings to a colleague who recorded the results. When performing repeat readings of multiple plates, the technologist cycled the different plates.

Caliper readings. Six technologists in the clinical microbiology laboratory participated in these studies. They placed the K-B plates on the viewer and measured zones by using a digital caliper (Sylvac; Fowler Tools and Instruments, Boston, Mass.) connected to a foot pedal. When the pedal was depressed, the zone diameter automatically entered a database (WHONET) on a desktop personal computer.

Sirscan readings. After putting the K-B plate on a sliding tray, a keystroke initiates the zone readings. An image of the plate appears on the screen, with the zone diameters encircled by a green (susceptible), yellow (intermediate), or red (resistant) circle. At this point the reader can modify the zone readings. Another keystroke automatically enters the values into the Sirscan database. The timing tests performed on the quality control strains and the three clinical isolates were done without modifying the Sirscan readings. In the prospective

^{*} Corresponding author. Mailing address: Miriam Hospital, 164 Summit Ave., Providence, RI 02906. Phone: (401) 793-4622. Fax: (401) 751-2398. E-mail: Antone_Medeiros@brown.edu.

Isolate, reader(s), and method $(n)^{\alpha}$	Time (s), (mean ± SD)					Diam (n	nm, mean ± S	D) of zone of	inhibition ^b				
<i>E. coli</i> (CI) J.C., ruler (5) S.M., caliper (5) 4 Techs, caliper (4) J.C., Sirscan (10)	$\begin{array}{c} 39.4 \pm 2.7 \\ 26.2 \pm 2.0 \\ 32 \pm 6.0 \\ 10.7 \pm 0.6 \end{array}$	FUR 19.8 ± 0.4 19.8 ± 0.4 19.8 ± 0.4 20 ± 0	$\begin{array}{c} {\rm KEF} \\ 17\pm 0 \\ 17\pm 0 \\ 17.3\pm 0.4 \\ 16.6\pm 0.5 \end{array}$	AMP 19 ± 0 19.2 ± 0.4 19.5 ± 0.5 19 ± 0	$\begin{array}{c} {\rm GEN}\\ 18\pm 0\\ 18\pm 0.6\\ 18.8\pm 1.3\\ 20.6\pm 0.7 \end{array}$	$SXT 25.8 \pm 0.4 25.6 \pm 0.5 26 \pm 0 27.7 \pm 0.6 $	FTX 28.8 \pm 0.4 28.8 \pm 0.4 29.8 \pm 0.8 31.8 \pm 0.4	$\begin{array}{c} \text{IMP} \\ \text{27.4} \pm 0.5 \\ \text{27.2} \pm 0.4 \\ \text{28.3} \pm 0.8 \\ \text{27.7} \pm 0.9 \end{array}$	AUG 19.6 \pm 0.5 19.4 \pm 0.5 19.8 \pm 0.8 19 \pm 0	FRX 22 ± 0.9 21.8 ± 0.4 22.3 ± 1.5 22.2 ± 0.4	MEZ 24 ± 0 24 ± 0 25 ± 0 25.2 ± 0.4	NOR 25.6 \pm 0.8 26 \pm 0.6 27.3 \pm 1.3 31.9 \pm 0.3	ATM 28.6 ± 0.5 28.4 ± 0.5 29 ± 1.6 31.2 ±
<i>M. morganii</i> (CI) J.C., ruler (5) S.M., caliper (5) 4 Techs, caliper (4) J.C., Sirscan (10)	$\begin{array}{c} 37.2 \pm 3.4 \\ 26.2 \pm 4.5 \\ 33 \pm 6.7 \\ 11 \pm 0.6 \end{array}$	KEF 6 ± 0 6 ± 0 6.3 ± 0.4 6 ± 0	AMP 6 ± 0 6 ± 0 6 ± 0 6 ± 0	GEN 17. ± 0 17 ± 0 17.3 ± 0.4 17.7 ± 1.0	$\begin{array}{c} \text{SXT} \\ \text{22.2 } \pm 0.7 \\ \text{22.2 } \pm 0.4 \\ \text{22.3 } \pm 1.3 \\ \text{25.7 } \pm 1.9 \end{array}$	FTX 17 ± 0 16.6 ± 0.5 17 ± 0 17 ± 0 17 ± 0	$\begin{array}{c} \text{IMP} \\ 19 \pm 0.6 \\ 19.2 \pm 0.4 \\ 19.5 \pm 0.5 \\ 23.3 \pm 0.5 \end{array}$	FRX 8 ± 0 6.4 ± 0.8 7.5 ± 0.9 6 ± 0	$\begin{array}{c} \text{MEZ} \\ 11.2 \pm 0.4 \\ 12.4 \pm 0.5 \\ 12 \pm 0.7 \\ 13.5 \pm 0.7 \end{array}$	$\begin{array}{c} \text{ATM} \\ 21.2 \pm 1.2 \\ 22 \pm 0 \\ 23 \pm 1.2 \\ 24.2 \pm 0.4 \end{array}$	FOX 16 ± 0 16 ± 0 16.5 ± 0.5 16.2 ± 0.4	$\begin{array}{c} \text{PIP} \\ 16 \pm 0.6 \\ 15.8 \pm 0.4 \\ 16 \pm 0.7 \\ 16 \pm 0 \end{array}$	CIP 28.4 ± 0.8 27.6 ± 0.5 28.8 ± 2.3 366 ± 2.2
S. <i>aureus</i> (CI) J.C., ruler (5) S.M., caliper (5) 4 Techs, caliper (4) J.C., Sirscan (10)	$\begin{array}{c} 27 \pm 2.8 \\ 17.6 \pm 1.2 \\ 21.8 \pm 5.4 \\ 11 \pm 0.6 \end{array}$	$\begin{array}{c} {\rm GEN}\\ 20.2\pm0.4\\ 20\pm0\\ 20.5\pm0.5\\ 20.8\pm0.4\end{array}$	$\begin{array}{c} {\rm AUG}\\ {\rm 26.6 \pm 0.5}\\ {\rm 26 \pm 0}\\ {\rm 26.8 \pm 0.8}\\ {\rm 27 \pm 0}\end{array}$	CIP 21 ± 0 21 ± 0 22 ± 1.2 21.4 ± 0.5	PEN 14.0 ± 0 14.2 ± 0.4 14.5 ± 0.5 14.6 ± 0.5	ERY 6 ± 0 6 ± 0 6.5 ± 0.9 9.6 ± 0.5	CLI 23.6 \pm 0.5 24 \pm 0 24 \pm 0.7 27.2 \pm 0.7	OXA 17.8 ± 0.4 18 ± 0 18.8 ± 0.4 17.4 ± 0.5	$VAN \\ 18.8 \pm 0.4 \\ 18.8 \pm 0.4 \\ 19.3 \pm 0.8 \\ 18 \pm 0.8 \\ 18 \pm 0$				
<i>E. coli</i> 25922 (QC) L.S., caliper (6) 3 Techs, caliper (3) J.C., Sirscan (10)	$\begin{array}{c} 22 \pm 2.6 \\ 29 \pm 5.1 \\ 11.1 \pm 0.3 \end{array}$	KEF 18.3 ± 0.5 17.3 ± 0.5 17.2 ± 0.4	AMP 19.7 ± 0.5 18.7 ± 0.9 18.2 ± 0.4	GEN 19 ± 0.6 19.3 ± 0.5 21.8 ± 0.9	$\begin{array}{c} \text{SXT} \\ 25 \pm 0.6 \\ 25 \pm 0 \\ 25 \pm 0 \end{array}$	FRX 29.3 \pm 0.9 29 \pm 0 31.9 \pm 0.3	$IMP25.7 \pm 0.726 \pm 0.830.3 \pm 0.5$	FRX 22.7 ± 0.5 21.7 ± 0.9 22.3 ± 0.5	MEZ 24.7 ± 0.5 23.3 ± 0.5 25.2 ± 0.4	ATM 29.17 \pm 1.1 29 \pm 0.8 31.9 \pm 0.3	FOX 25.3 ± 0.5 25.3 ± 0.5 24.8 ± 0.6	PIP 25 ± 0 25 ± 0.8 27.2 ± 0.4	CIP 31.5 \pm 1.3 31.7 \pm 0.5 36 \pm 0
P. aeruginosa 27853 (QC) L.S., caliper (6) 3 Techs, caliper (3) J.C., Sirscan (10)	$\begin{array}{c} 20.5 \pm 1.0 \\ 29.7 \pm 5.1 \\ 11 \pm 0 \end{array}$	GEN 18.3 ± 0.5 18 ± 0 18.1 ± 0.3	$\begin{array}{c} \text{SXT} \\ 6 \pm 0 \\ 6 \pm 0 \\ 6 \pm 0 \end{array}$	$IMP23 \pm 0.622.3 \pm 0.523 \pm 0.6$	$\begin{array}{c} \text{MEZ} \\ 19.8 \pm 0.7 \\ 20 \pm 0 \\ 20.1 \pm 0.3 \end{array}$	ATM 25.3 ± 0.5 25 ± 0.8 26 ± 0	PIP 26.2 ± 0.4 26.3 ± 0.5 28.9 ± 0.3	CIP 25.5 \pm 0.5 26 \pm 1.4 30.8 \pm 0.6	CHL 6 ± 0 6 ± 0 6 ± 0	TOB 20.8 ± 0.4 20.3 ± 0.5 20.3 ± 0.6	AMK 21.67 ± 0.5 21.7 ± 0.5 21.8 ± 0.9	$\begin{array}{c} \text{CAR} \\ 19.5 \pm 0.8 \\ 20 \pm 0 \\ 21.1 \pm 0.3 \end{array}$	CAZ 26.3 ± 0.5 26 ± 1.4 27 ± 0
S. aureus 25923 (QC) L.S., caliper (6) 3 Techs, caliper (3) J.C., Sirscan (10)	$\begin{array}{c} 15.5 \pm 1.3 \\ 21 \pm 2.4 \\ 11 \pm 0 \end{array}$	GEN 22.5 ± 0.5 20.6 ± 0.5 20.9 ± 0.3	$\begin{array}{c} \text{SXT} \\ 27.7 \pm 0.5 \\ 27.7 \pm 0.5 \\ 29.6 \pm 0.5 \end{array}$	AUG 30.8 ± 0.4 30.3 ± 1.2 32.5 ± 0.5	CIP 23.5 \pm 0.8 23.7 \pm 0.9 25 \pm 0	PEN 34.7 ± 0.5 33.7 ± 0.9 34 ± 0	ERY 26.2 ± 0.4 24.7 ± 0.9 27.1 ± 0.3	CLI 25.7 ± 0.5 24.3 ± 0.5 28.1 ± 0.5	EXA 24.2 ± 0.4 23 ± 0.8 24.4 ± 0.5	VAN 18.8 \pm 0.4 17.7 \pm 0.5 17 \pm 0			
^a Readers are indicated by i ^b FUR, nitrofurantoin; KEI mezlocillin; NOR, norfloxacin;	initials. Techs, t F, cephalothin; ATM, aztreona	chnicians; CI, o AMP, ampicil am; FOX, cefo	clinical isolate; lin; GEN, gen xitin; PIP, pipe	QC, quality c tamicin; SXT; racillin; CIP, c	ontrol strain; , , trimethoprin siprofloxacin; I	<i>n</i> , number of 1 n-sulfamethoxa PEN, penicillir	reads. nzole; FTX, co 1; ERY, erythr	efotaxime; IM omycin; CLI, c	P, imipenem; lindamycin; O	AUG, amoxicil XA, oxacillin; V	lin-clavulanate; ⁷ AN, vancomyc	; FRX, cefuro sin; CHL, chlor	xime; MEZ, amphenicol;

meziociinii, NOK, noritoxacii, ATM, aztreonani, FOA, cetoxittii, FTF, piperaciinii, TOB, tobramycin; AMK, amikacin; CAR, carbenicillin; CAZ, ceftazidime.

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TABLE 1. Repeated zone diameter measurements of three clinical isolates and three quality control strains by ruler, caliper, and Sirscan



FIG. 1. Correlation of zone diameters as measured with a Sirscan image analyzer to caliper readings. Dark lines represent the NCCLS breakpoints. \bigcirc , automatic reading; X, reviewed reading. (A) Penicillin disk with 127 clinical isolates of staphylococci. (B) Vancomycin disk with 127 staphylococci. (C) Ampicillin disk with 209 gram-negative isolates.

study of 368 clinical isolates, the reader did not adjust the Sirscan readings (automatic readings) with the first 114 isolates; the remainder were adjusted as judgment dictated (reviewed readings).

Timing studies. A stopwatch was used to time all readings. Timing began when the technologist commenced reading the zone sizes of the K-B plate on the viewer. With the Sirscan, timing began when the keystroke initiated the reading of the K-B plate in the loading tray.

Time required to measure zone diameters. Table 1 shows the results of readings of clinical isolates of *S. aureus, E. coli*, and *Morganella morganii* by the three different methods. The median time to read the 12 disk diameters on the plates with gram-negative bacilli was 39 s by ruler, 28 s by caliper, and 11 seconds by Sirscan.

Reading the eight disks on the *S. aureus* plate took 27 s by ruler, 18 s by caliper, and 11 s by Sirscan. The reading times of four different technologists using the caliper ranged from 24 to 41 s for the *E. coli* plates, 22 to 44 s for the *M. morganii* plates, and 17 to 31 s for the *S. aureus* plates. Sirscan reading times varied between 10 and 12 s. Similar results were observed with the control strains.

Zone size variation upon repeated testing. Zone sizes of control strains varied by more than 2 mm in 1 of 340 (0.3%) determinations by Sirscan and 6 of 297 (2.0%) readings by caliper. With three clinical isolates, zone diameters varied more than 2 mm in 4 of 297 (1.3%) readings by Sirscan and 7 of 330 (2.1%) readings by caliper. Variation exceeded 3 mm in 4 of 1,254 (0.3%) readings.

Comparison of zone sizes measured by a Sirscan automated reader with those measured by caliper. Over a 4-month period, 368 fresh clinical isolates were tested prospectively by both methods in the Clinical Microbiology Laboratory. Correlation of zone diameter sizes was very good with penicillin, vancomycin, and ampicillin disks (Fig. 1). There was also good correlation of diameters for oxacillin and amoxicillin-clavulanate, although Sirscan failed to detect light growth around the oxacillin disk in 3 of 131 (2.3%) isolates and around the amoxicillin-clavulanate disk in 2 of 219 isolates (0.9%) (Fig. 2). Correlation of zone sizes was good, but not as tight, for imipenem and cephalothin, perhaps due to species-specific variation in growth around these disks (Fig. 3). In this regard, zone diameters around imipenem disks were 2 to 3 mm larger by Sirscan than by caliper with the control clinical isolates of M. morganii, but not with the E. coli or P. aeruginosa isolates.

Sirscan zone diameter measurements tended to be 3 to 5 mm larger than caliper readings around ciprofloxacin, norfloxacin, aztreonam, trimethoprim-sulfamethoxazole, and nitrofurantoin disks when testing gram-negative bacilli (Fig. 4) and around erythromycin and clindamycin disks with staphylococci (Fig. 5). The effect tended to be more pronounced the larger the zone diameter, especially with nitrofurantoin. A less pronounced shift of zone sizes, about 2 to 3 mm greater with Sirscan than by caliper, occurred with piperacillin, mezlocillin,



FIG. 2. Correlation of zone diameters as measured with a Sirscan image analyzer to caliper readings. O, automatic reading; X, reviewed reading. (A) Oxacillin disk with 53 clinical isolates of *S. aureus* and 74 coagulase-negative staphylococci. (B) Amoxicillin-clavulanate disk with 127 staphylococci and 87 gram-negative bacilli. The NCCLS breakpoints are not shown because they differ for the different groups of bacteria.



FIG. 3. Correlation of zone diameters as measured with a Sirscan image analyzer to caliper readings. Dark lines represent the NCCLS breakpoints. O, automatic reading; X, reviewed reading. (A) Imipenem disk with 241 gram-negative isolates. (B) Cephalothin disk with 210 gram-negative isolates.

cefuroxime, cefotaxime, and gentamicin (Fig. 6). There was little difference in these shifts between the automatic and reviewed readings.

Very major errors (resistant by caliper and susceptible by Sirscan) occurred with 10 of 3,770 (0.3%) readings (Table 2). Eight of the 10 very major errors were with antibiotic disks that had no intermediate zone separating the resistant and susceptible populations. They occurred in testing staphylococci with amoxicillin-clavulanate (5 of 128, 3.9%), pseudomonas with piperacillin (1 of 28, 3.6%), coagulase-negative staphylococci with oxacillin (2 of 74, 2.7%), gram-negative bacilli with cefuroxime (1 of 209, 0.5%), and mixed species with trimethoprim-sulfamethoxazole (1 of 366, 0.3%). The percentages of results that were very major errors were not statistically different between the automatic and reviewed readings.

Discussion. The Sirscan image analyzer reads zone diameters more than twice as rapidly as skilled technologists using a computer interfaced caliper system. Reproducibility of measurements was excellent, within about 1 mm for Sirscan and 3 mm for caliper readings. The latter, however, tended to vary with the technologist, unlike readings made with the Sirscan.

With regard to concordance of zone diameter measurements, the Sirscan often failed to detect light growth at the margins ("beach effect") of the inhibition zones of some disks. Consequently, zone diameters measured by Sirscan were significantly larger for ciprofloxacin, norfloxacin, aztreonam, erythromycin, clindamycin, and trimethoprim-sulfamethoxazole. However, these differences rarely affected the classification of the isolate as susceptible or resistant. The overall frequency of very major errors (resistant by caliper and susceptible by Sirscan) was low, i.e., 0.3% (10/3,770 readings). Three of these errors occurred with the oxacillin disk, reflecting the difficulty of visualizing the light growth that sometimes occurs around the oxacillin disk with resistant *S. aureus* organisms. Another possible source of error is failure to swab the K-B plate thoroughly. If growth is not confluent, the Sirscan



FIG. 4. Correlation of zone diameters as measured with a Sirscan image analyzer to caliper readings. Dark lines represent the NCCLS breakpoints. O, automatic reading; X, reviewed reading. (A) Ciprofloxacin disk with 278 staphylococci and gram-negative bacilli. (B) Aztreonam disk with 241 gram-negative isolates. (C) Trimethoprim-sulfamethoxazole disk with 366 staphylococci and gram-negative bacilli. (D) Nitrofurantoin disk with 115 staphylococci and gram-negative bacilli. Norfloxacin readings are not shown.

Antibiotic(s)	Isolate(s)	NCCLS breakpoints (mm)	No. tested	SIc/SIs ^a	Rc/Rs ^b	SIc/Rs ^c	Rc/SIs ^d	Zone size(s) (mm) of Rc/SIs isolate(s) ^e	% of results that were very major errors
Amoxicillin- clavulanate	Staphylococcus sp.	≥20	127	102	19	1	5	20 (S), 20 (S), 20 (S), 21 (S), 22 (S)	3.9
	Gram-negative bacilli	14–17	87	73	14	0	0		
Ampicillin	Gram-negative bacilli	14-16	209	87	122	0	0		
Aztreonam	Gram-negative bacilli	16–21	241	219	18	0	4	16 (I), 17 (I), 17 (I), 17 (I)	
Cefotaxime	Gram-negative bacilli	15-22	209	197	10	1	1	15 (I)	
Cefuroxime sodium	Gram-negative bacilli	15-17	209	171	37	0	1	18 (S)	0.5
Cephalothin	Gram-negative bacilli	15-17	210	128	75	7	0		
Ciprofloxacin	Gram-negative bacilli and Staphylococcus sp.	16-20	278	221	56	0	1	17 (I)	
Clindamycin	Staphylococcus sp.	15 - 20	127	99	28	0	0		
Erythromycin	Staphylococcus sp.	14-22	127	46	79	0	2	14 (I), 14 (I)	
Gentamicin	Gram-negative bacilli and Staphylococcus sp.	13–14	364	326	35	1	2	14 (I), 14 (I)	
Imipenem	Gram-negative bacilli	14-15	241	233	8	0	0		
Mezlocillin	Pseudomonas sp.	≥16	28	18	10	0	0		
	Other gram-negative bacilli	18-20	212	159	48	4	1	18 (I)	
Nitrofurantoin	Gram-negative bacilli and Staphylococcus sp.	15–16	115	92	18	0	5	15 (I), 15 (I), 15 (I), 15 (I), 15 (I)	
Norfloxacin	Gram-negative bacilli	18 - 20	87	1	86	0	0		
Oxacillin	Coagulase-negative staphylococci	≥ 18	74	19	53	0	2	22 (S), 25 (S)	2.7
	S. aureus	11-12	53	38	15	0	0		
Penicillin G	Staphylococcus sp.	28-29	127	115	12	0	0		
Piperacillin	Pseudomonas sp.	≥ 18	28	26	1	0	1	19 (S)	3.6
1	Other gram-negative bacilli	18-20	124	92	31	1	0		
Trimethoprim- sulfamethoxazole	Gram-negative bacilli and Staphylococcus sp.	11–15	366	266	96	2	2	16 (S), 12 (I)	0.3
Vancomycin	Staphylococcus sp.	≥15	127	127	0	0	0		

TABLE 2. Isolates susceptible, intermediate, or resistant as determined by caliper and Sirscan readings of zone diameters

^a Number of isolates susceptible or intermediate by both caliper and Sirscan.

^b Number of isolates resistant by both caliper and Sirscan.

^c Number of isolates susceptible or intermediate by caliper and resistant by Sirscan.

^d Number of isolates resistant by caliper and susceptible or intermediate by Sirscan.

^e S, sensitive; I, intermediate; R, resistant.

f (Number of Rc/SIs isolates) – (number of intermediate isolates)/number of isolates tested \times 100.

may read between the growth. A technician screening the Sirscan readings prior to validation and modifying them as needed can minimize these errors.

A study comparing the Sirscan readings to manual readings done in four laboratories in France reported a higher rate (1.76%) of very major errors, which varied according to species, antibiotic, and hospital (1). *Burkholderia cepacia*, *Staphylococcus epidermidis*, *Stenotrophomonas maltophilia*, and *Branhamella* isolates were especially problematical. Differences between the two studies are probably due to the mix of species and antibiotics tested. For example, our study did not include *Branhamella* isolates and the antibiotics fosfomycin and cefoperazone, which had high discordance rates. Interestingly, their study showed that square petri dishes yielded more reliable results than round dishes.

The Sirscan image analyzer merits strong consideration as a method of measuring and recording zone diameters. It provides a fully quantitative measure of antibiotic resistance, an important parameter for tracking emerging mechanisms of resistance and their spread in hospital bacteria (2). The computer interface with an expert system that screens each result according to user-defined algorithms enhances quality control. It also would make it feasible to employ species-specific breakpoints if and when such criteria are developed (3, 5). A limi-



FIG. 5. Correlation of zone diameters as measured with a Sirscan image analyzer to caliper readings. Dark lines represent the NCCLS breakpoints. \bigcirc , automatic reading; X, reviewed reading. (A) Erythromycin disk with 127 clinical isolates of staphylococci. (B) Clindamycin disk with 127 clinical isolates of staphylococci.



FIG. 6. Correlation of zone diameters as measured with a Sirscan image analyzer to caliper readings. Dark lines represent the NCCLS breakpoints. \bigcirc , automatic reading; X, reviewed reading. (A) Piperacillin disk with 28 *Pseudomonas* and 124 other gram-negative isolates. (B) Cefuroxime disk with 209 gram-negative isolates. (C) Cefotaxime disk with 209 gram-negative isolates. (D) Gentamicin disk with 364 isolates of staphylococci and gram-negative bacilli. Mezlocillin readings are not shown.

tation of the Sirscan is its inability to read plates inoculated with enterococcus or haemophilus species due to their light growth.

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