Comparison of MIC Results for Gepotidacin by Agar Dilution and Broth Microdilution Methods for Various Gram-negative and Gram-positive Species

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Introduction

- Gepotidacin (GSK2140944) is a novel, bactericidal, first in class triazaacenaphthylene antibiotic in clinical development for the treatment of gonorrhea and uncomplicated urinary tract infection (acute cystitis).
- Gepotidacin selectively inhibits bacterial DNA replication by a distinct mechanism of action which confers in vitro activity against most strains of target pathogens, such as E. coli, S. saprophyticus and N. gonorrhoeae, including those resistant to current antibiotics.
- The aim of this study, in accordance with Clinical and Laboratory Standards Institute (CLSI) M23-A4 guidelines (2018), was to test the equivalency of minimal inhibitory concentrations (MICs) obtained by 2 reference antimicrobial susceptibility testing methods, agar dilution and broth microdilution, for the investigational antimicrobial drug gepotidacin against various gram-positive and gram-negative organisms.

Materials and Methods

- Susceptibility testing for both methods was performed for 857 clinical isolates recovered from over 120 medical centers worldwide.
- All isolates were collected in 2020, except for 28 *Providencia rettgeri* isolates from 2019 due to lower prevalence.
- MICs were determined by broth microdilution and agar dilution per CLSI M07-A10 methods (2018).
- Cation-adjusted Mueller-Hinton broth or Mueller-Hinton agar (MHA) was used for all organisms.
- Broth microdilution and agar dilution for all isolates were performed on the same day from the same bacterial inoculum suspension.
- Due to the swarming nature of many Proteus mirabilis isolates, testing was carried out in 24-well, non-treated microtiter plates. A total volume of 600 µL of molten agar with the various drug concentrations was transferred into each well. After the agar was allowed to cool, each well was inoculated with a single isolate suspension at the appropriate concentration.
- Quality control (QC) strains were tested concomitantly with clinical isolates, and inoculum density was monitored by colony counts for the following QC strains: Escherichia coli ATCC 25922, Enterococcus faecalis ATCC 29212, Pseudomonas aeruginosa ATCC 27853, and Staphylococcus aureus ATCC 29213.
- Four ATCC QC isolates were evaluated with at least 21 replicate MIC results from at least 2 testing days and analyzed for intra- and interday reproducibility, including range, mean, mode, and standard deviation.
- Essential agreement (EA) based on evaluable results was calculated as the number of isolates with MICs within one 2-fold dilution of the reference method divided by the total number of results.
- Equivalency was defined using the 95% criteria from the Food and Drug Administration's class II controls document (2009).

Results

- Replicate testing against QC organisms demonstrated limited variability. No drug/isolate/MIC method combination had MIC values that differed by more than 1-log, dilution from the mode (Table 1).
- The modal gepotidacin MIC value for *E. faecalis* ATCC 29212 was 2-fold higher when determined by broth microdilution (2 µg/mL) than by agar dilution (1 µg/mL). Similarly, the modal gepotidacin MIC value for S. aureus ATCC 29213 was 2-fold higher when determined by broth microdilution (0.25 µg/mL) than by agar dilution (0.12 μ g/mL; Table 1).
- When comparing all isolates, gepotidacin MIC values by these 2 methods showed good correlation $(R^2 = 0.91)$ and an EA of 92.5% (Figure 1 and Table 2).
- For most species, broth microdilution MICs were often 1-log, dilution higher than agar dilution MICs (Table 3).
- Against *E. faecalis*, broth microdilution MIC_{50/90} 2/2 μ g/mL and agar dilution MIC_{50/90} 1/1 μ g/mL were observed. The trend of higher MIC values determined by broth microdilution led to an EA of only 84.1% with the agar dilution method.

- Among Staphylococcus saprophyticus isolates, broth microdilution gepotidacin MICs were similar to agar dilution MICs (MIC_{50/90}; 0.06/0.12 µg/mL, 0.06/0.06 µg/mL respectively). The EA for S. saprophyticus was 90.7%.
- Against *Klebsiella* species isolates tested, EAs observed were 92.2% for all isolates, 95.4% for K. pneumoniae, 98.1% for K. aerogenes, but only 80.0% for K. oxytoca.
- For Enterobacter cloacae complex species and Citrobacter species, broth microdilution MIC values were often 1-log, dilution higher than agar dilution MIC values. However, EAs observed were 95.3% for *E. cloacae* complex species, 94.3% for *C. freundii*, and 100.0% for *C. koseri*.
- Against *P. rettgeri*, broth microdilution MIC_{50/90} 8/16 μ g/mL and agar dilution MIC_{50/90} 4/8 μ g/mL were observed. The trend of higher MIC values determined by broth microdilution led to an EA of only 88.6% with the agar dilution method.
- Against *P. mirabilis*, broth microdilution gepotidacin MIC_{50/90} values were equivalent to agar dilution MIC values for this species (MIC_{50/90}; 8/16 µg/mL) and all MIC values (100%) showed EA between broth microdilution and agar dilution.

Conclusions

- Limited intra- or interday variation was observed for MICs against the 4 QC strains and no isolate/testing method combination had MIC values that differed by more than 1-log, dilution from the mode.
- An EA between broth microdilution and agar dilution methods of 92.5% was observed for all isolates tested.
- Slightly higher gepotidacin MICs, in most cases one 2-fold dilution higher, were observed when tested by broth microdilution for most species/groups tested.
- With an EA of >95%, equivalency was established between agar dilution and broth microdilution MIC methods for gepotidacin against K. pneumoniae, K. aerogenes, E. cloacae species complex, P. mirabilis, and C. koseri.
- Equivalency was not established between methods for *C. freundii* species complex, *S. saprophyticus*, P. rettgeri, E. faecalis, and K. oxytoca.

Table 1. Gepotidacin broth microdilution and agar dilution MIC results when tested against quality control strains *E. coli* ATCC 25922, *E. faecalis* ATCC 29212, P. aeruginosa ATCC 27853, and S. aureus ATCC 29213

Organism (no. of tests / no. of days)	MIC (µg/mL)									
Antimicrobial agent and testing method	Range	Mean	Standard Deviation	Mode						
E. coli ATCC 25922										
Gepotidacin agar dilution (21/9) ^a	1	1	0	1						
Gepotidacin broth microdilution (24/12)	1 – 2	1.1	0.34	1						
E. faecalis ATCC 29212										
Gepotidacin agar dilution (21/9)	1	1	0	1						
Gepotidacin broth microdilution (24/12)	1 – 4	2	0.51	2						
P. aeruginosa ATCC 27853										
Gepotidacin agar dilution (21/9)	8	8	0	8						
Gepotidacin broth microdilution (24/12)	4 – 16	8.3	2.62	8						
S. aureus ATCC 29213										
Gepotidacin agar dilution (21/9)	0.12 – 0.25	0.13	0.04	0.12						
Gepotidacin broth microdilution (24/12)	0.25	0.25	0	0.25						

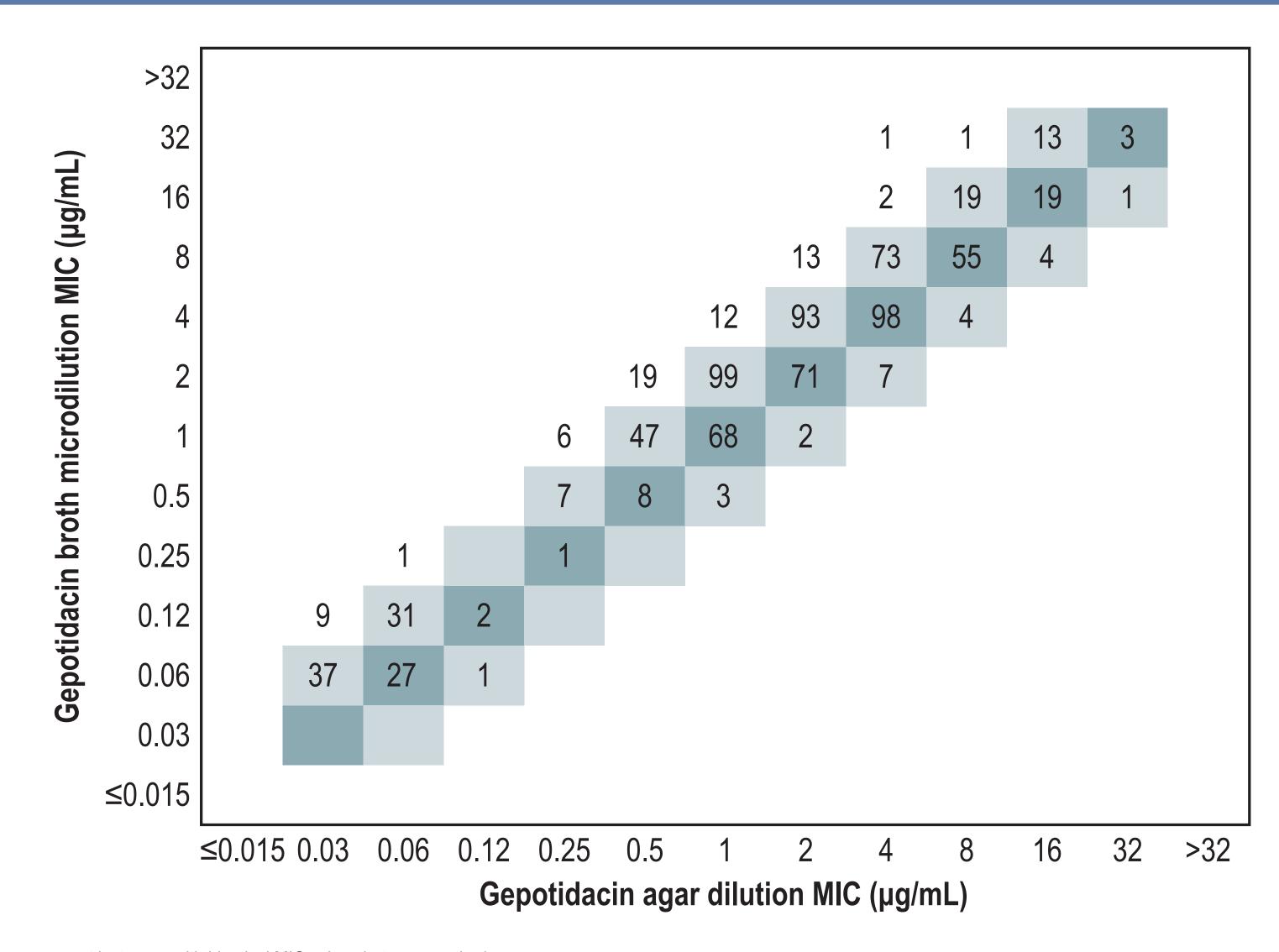
^a MIC results from 24-well agar *P. mirabilis* testing not included in this analysis

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			ion differe	ence) ^a						
	% Essential agreement ^b	← Larger agar dilution MIC			Larger broth microdilution MIC \rightarrow					
Organism/organism group		≤-3	-2	-1	0	1	2	≥3		
All isolates	92.5%			22	352	419	63	1		
Enterococcus faecalis	84.1%			1	20	69	17			
Staphylococcus saprophyticus	90.7%			1	29	68	10			
Klebsiella species	92.2%			1	93	107	17			
Klebsiella pneumoniae	95.4% ^c			1	45	58	5			
Klebsiella aerogenes	98.1%				30	23	1			
Klebsiella oxytoca	80.0%				18	26	11			
Enterobacter cloacae species complex	x 95.3%			2	52	48	5			
Providencia rettgeri	88.6%			3	29	61	11	1		
Proteus mirabilis	100.0%			11	73	20				
Citrobacter species	97.2%			3	56	46	3			
Citrobacter freundii species complex	94.3%			2	30	18	3			
Citrobacter koseri	100.0%			1	26	28				

The log₂ dilution difference calculated as the log₂ (broth microdilution MIC value/agar dilution MIC value). Essential agreement calculated as the number of broth microdilution and agar dilution MIC values for each isolate that are ±1 log₂ dilution (shaded values), divided by the total number of isolates. Bold values denote essential agreement values >95%

Figure 1. Scattergram of gepotidacin broth microdilution MIC vs gepotidacin agar dilution MIC against all isolates (n=857) with an essential agreement of 92.5%



Dark blue shaded values represent instances with identical MIC values between methods ight blue shaded values represent instances of agreement included in essential agreement calculations (±1 log, dilution).

Acknowledgements/Disclosures

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Table 3. Antimicrobial activity of gepotidacin tested by broth microdilution and agar dilution against the clinical isolates tested														
Organism/test method		No. and cumulative % of isolates at MIC (µg/mL) of :												
(no. of isolates)	≤0.015						1	2	4	8	16	>16	MIC ₅₀	
Enterococcus faecalis (107)				0	1	1	30	60	3	1				
Broth microdilution method				0 0.0	0.9	4.7	32.7	68 96.3	-	100.0			2	2
Agar dilution method				0 0.0	6 5.6	31 34.6	60 90.7	8 98.1	2 100.0				1	1
Staphylococcus saprophyticus (108)														
Broth microdilution method		0 0.0	65 60.2	42 99.1	1 100.0								0.06	0.12
Agar dilution method	0 0.0	46 42.6	59 97.2	3 100.0									0.06	0.06
Klebsiella pneumoniae (109)														
Broth microdilution method					0 0.0	1 0.9	1 1.8	5 6.4	62 63.3	23 84.4	8 91.7	9 100.0	4	16
Agar dilution method				0 0.0	1 0.9	1 1.8	3 4.6	31 33.0	51 79.8	11 89.9	11 100.0		4	16
Klebsiella aerogenes (54)				0.0	0.3	1.0	4.0	55.0	73.0	03.3	100.0			
Broth microdilution method						0 0.0	6 11.1	38 81.5	10 100.0				2	4
Agar dilution method				0 0.0	1 1.9	0	20 38.9	31 96.3	2 100.0				2	2
Klebsiella oxytoca (55)				0.0	1.0	1.0								
Broth microdilution method						0 0.0	18 32.7	25 78.2	10 96.4	2 100.0			2	4
Agar dilution method				0 0.0	1 1.8	4 9.1	43 87.3	5 96.4	2 100.0				1	2
<i>Enterobacter cloacae</i> species complex (107)														
Broth microdilution method							0 0.0	29 27.1	53 76.6	12 87.9	9 96.3	4 100.0	4	16
Agar dilution method						0 0.0	7 6.5	53 56.1	32 86.0	8 93.5	5 98.1	2 100.0	2	8
Providencia rettgeri (105)														
Broth microdilution method						0.0	4 3.8	/ 10.5	36 44.8	47 89.5	7 96.2	4 100.0	8	16
Agar dilution method					0 0.0	3 2.9	5 7.6	25 31.4	56 84.8	13 97.1	3 100.0		4	8
Proteus mirabilis (104)														
Broth microdilution method					0 0.0	1 1.0	4 4.8	5 9.6	21 29.8	56 83.7	16 99.0	1 100.0	8	16
Agar dilution method						0 0.0	4 3.8	9 12.5	28 39.4	45 82.7	16 98.1	2 100.0	8	16
<i>Citrobacter freundii</i> species complex (53)														
Broth microdilution method					0	5 9.4	24 54.7	14 81.1	8 96.2	1 98.1	1 100.0		1	4
Agar dilution method				0 0.0	2 3.8	14 30.2	18 64.2	13 88.7	4 96.2	1 98.1	1 100.0		1	4
Citrobacter koseri (55)											10010			
Broth microdilution method					0	7 12.7	36 78.2	5 87.3		3 100.0			1	4
Agar dilution method				0 0.0	3 5.5	21 43.6	22 83.6	4 90.9	4 98.2	1 100.0			1	2

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