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- The present **MIC Test Strip Interpretative Criteria and Quality Control** table does not replace the official documents by CLSI and EUCAST.
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ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL
AMIKACIN 0.016 - 256 µg/mL	AK	Enterobacteriaceae <i>P. aeruginosa</i> <i>Acinetobacter</i> spp. Other Non-Enterobacteriaceae <i>Staphylococcus</i> spp.	16 16 16 16 16		32 32 32 32 32	64 64 64 64 64	Enterobacteriaceae <i>Pseudomonas</i> spp. <i>Acinetobacter</i> spp. <i>S. aureus</i> Coagulase-negative Staphylococci PK/PD (Non-species related) breakpoints	8 8 8 8 8 8	16 16 16 16 16 16	<i>S. aureus</i> ATCC® 29213 <i>E. faecalis</i> ATCC® 29212 <i>E. coli</i> ATCC® 25922 <i>P. aeruginosa</i> ATCC® 27853	1-4 64-256 0.5-4 1-4
AMOXICILLIN 0.016 - 256 µg/mL	AML	<i>S. pneumoniae</i> (nonmeningitis)	2		4	8	Enterobacteriaceae <i>Enterococcus</i> spp. Viridans group streptococci <i>H. influenzae</i> <i>N. meningitidis</i> Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes <i>H. pylori</i> <i>P. multocida</i> <i>Kingella kingae</i> PK/PD (Non-species related) breakpoints	8 4 0.5 2 0.12 4 0.5 0.12 1 0.125 2	8 8 2 2 1 8 2 0.12 1 0.125 8	<i>K. pneumoniae</i> ATCC® 700603 <i>S. pneumoniae</i> ATCC® 49619 <i>E. coli</i> ATCC® 25922 <i>H. influenzae</i> ATCC® 49766	> 128 0.03-0.12 2-8 0.125-0.5
AMOXICILLIN* - CLAVULANIC ACID (2/1) 0.016 - 256 * µg/mL <i>CLSI recommended</i>	AUG	Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Haemophilus</i> spp. <i>S. pneumoniae</i> (nonmeningitis) Anaerobes	8 4 4 2 4		16 - - 4 8	32 8 8 8 16				<i>S. aureus</i> ATCC® 29213 <i>E. faecalis</i> ATCC® 29212 <i>E. coli</i> ATCC® 25922 <i>E. coli</i> ATCC® 35218 <i>K. pneumoniae</i> ATCC® 700603 <i>H. influenzae</i> ATCC® 49247 <i>S. pneumoniae</i> ATCC® 49619 <i>B. fragilis</i> ATCC® 25285 <i>B. thetaiotaomicron</i> ATCC® 29741	0.12-0.5 0.25-1 2-8 4-16 4-16 2-16 0.03-0.12 0.25-1 0.25-1
AMOXICILLIN* - CLAVULANIC ACID (2 µg/mL) 0.016 - 256 * µg/mL <i>EUCAST recommended</i>	AMC						Enterobacteriaceae Enterobacteriaceae (uncomplicated UTI only) <i>Enterococcus</i> spp. <i>H. influenzae</i> <i>M. catarrhalis</i> Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes <i>P. multocida</i> PK/PD (Non-species related) breakpoints	8 32 4 2 1 4 4 1 2	8 32 8 2 1 8 8 1 8	<i>E. coli</i> ATCC® 25922 <i>H. influenzae</i> ATCC® 49766 <i>E. coli</i> ATCC® 35218	2-8 0.125-0.5 4-32

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AMPICILLIN 0.016 - 256 µg/mL	AMP	Enterobacteriaceae Staphylococcus spp. Enterococcus spp. Haemophilus spp. Streptococcus spp. β-Hemolytic Group Streptococcus spp. Viridans Group N. meningitidis Anaerobes	8 0.25 8 1 0.25 0.25 0.12 0.5		16 - 2 - - 0.5-4 1	32 0.5 16 4 - 8 2 2	Enterobacteriaceae Enterococcus spp. S. pneumoniae Viridans group streptococci H. influenzae N. meningitidis Gram-positive anaerobes except C. difficile Gram-negative anaerobes L. monocytogenes P. multocida Aerococcus sanguinicola and urinae Kingella kingae PK/PD (Non-species related) breakpoints	8 4 0.5 0.5 1 0.12 4 0.5 1 1 1 0.25 0.06 2	8 8 2 2 1 1 8 2 1 1 0.25 0.06 8	S. aureus E. faecalis E. coli E. coli K. pneumoniae H. influenzae S. pneumoniae H. influenzae	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 35218 ATCC® 700603 ATCC® 49247 ATCC® 49619 ATCC® 49766	0.5-2 0.5-2 2-8 > 32 > 128 2-8 0.06-0.25 0.06-0.25
AMPICILLIN* - SULBACTAM (2/1) 0.016 - 256 * µg/mL CLSI recommended	AMS	Enterobacteriaceae Acinetobacter spp. Staphylococcus spp. Haemophilus spp. Anaerobes	8 8 8 2 8		16 16 16 - 16	32 32 32 4 32				E. coli E. coli K. pneumoniae H. influenzae B. fragilis B. thetaiotaomicron E. lenta	ATCC® 25922 ATCC® 35218 ATCC® 700603 ATCC® 49247 ATCC® 25285 ATCC® 29741 ATCC® 43055	2-8 8-32 8-32 2-8 0.5-2 0.5-2 0.5-2
AMPICILLIN* - SULBACTAM (4 µg/mL) 0.016 - 256 * µg/mL EUCAST recommended	SAM						Enterobacteriaceae Enterococcus spp. H. influenzae M. catarrhalis Gram-positive anaerobes except C. difficile Gram-negative anaerobes PK/PD (Non-species related) breakpoints	8 4 1 1 4 4 2	8 8 1 1 8 8 8	E. coli H. influenzae E. coli	ATCC® 25922 ATCC® 49766 ATCC® 35218	1-4 0.06-0.25 16-128
AZITHROMYCIN 0.016 - 256 µg/mL	AZM	S. Typhi Staphylococcus spp. H. influenzae S. pneumoniae Streptococcus spp. β-Hemolytic Group Streptococcus spp. Viridans Group N. meningitidis	16 2 4 0.5 0.5 0.5 2		- 4 - 1 1 1 -	32 8 - 2 2 2 -	Staphylococcus spp. Streptococcus groups A, B, C and G S. pneumoniae H. influenzae M. catarrhalis N. gonorrhoeae Kingella kingae	1 0.25 0.25 0.12 0.25 0.25 0.25	2 0.5 0.5 4 0.5 0.5 0.25	S. aureus H. influenzae (-CO <sub>2</sub> ) H. influenzae (+CO <sub>2</sub> ) S. pneumoniae (-CO <sub>2</sub> ) S. pneumoniae (+CO <sub>2</sub> ) N. gonorrhoeae	ATCC® 29213 ATCC® 49247 ATCC® 49247 ATCC® 49619 ATCC® 49619 ATCC® 49226	0.5-2 1-4 4-16 0.06-0.25 0.5-2 0.25-1
AZTREONAM 0.016 - 256 µg/mL 0.064 - 1024 µg/mL	ATM	Enterobacteriaceae P. aeruginosa Other Non-Enterobacteriaceae Haemophilus spp.	4 8 8 2		8 16 16 -	16 32 32 -	Enterobacteriaceae Pseudomonas spp. PK/PD (Non-species related) breakpoints	1 1 4	4 16 8	E. coli P. aeruginosa E. coli H. influenzae	ATCC® 25922 ATCC® 27853 ATCC® 35218 ATCC® 49247	0.06-0.25 2-8 0.03-0.12 0.12-0.5
CEFAZOLIN 0.016 - 256 µg/mL	CEC	Enterobacteriaceae Staphylococcus spp. Haemophilus spp. S. pneumoniae	8 8 8 1		16 16 16 2	32 32 32 4	S. pneumoniae H. influenzae	0.03 0.5	0.5 0.5	S. aureus E. coli H. influenzae S. pneumoniae	ATCC® 29213 ATCC® 25922 ATCC® 49766 ATCC® 49619	1-4 1-4 1-4 1-4
CEFAZOLIN 0.016 - 256 µg/mL	KZ	Enterobacteriaceae Enterobacteriaceae (surrogate test for uncomplicated UTI)	2 16		4 -	8 32	Viridans group streptococci PK/PD (Non-species related) breakpoints	0.5 1	0.5 2	S. aureus E. coli	ATCC® 29213 ATCC® 25922	0.25-1 1-4

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CEFEPIME 0.016 - 256 µg/mL 0.002 - 32 µg/mL	FEP	Enterobacteriaceae	2	4-8	-	16	Enterobacteriaceae	1	4	<i>S. aureus</i>	ATCC® 29213	1-4
		<i>P. aeruginosa</i>	8	-	16	32	<i>Pseudomonas</i> spp.	8	8	<i>E. coli</i>	ATCC® 25922	0.016-0.12
		<i>Acinetobacter</i> spp.	8	-	16	32	<i>S. pneumoniae</i>	1	2	<i>P. aeruginosa</i>	ATCC® 27853	0.5-4
		Other non-Enterobacteriaceae	8	-	16	32	Viridans group streptococci	0.5	0.5	<i>H. influenzae</i>	ATCC® 49247	0.5-2
		<i>Haemophilus</i> spp.	2	-	-	-	<i>H. influenzae</i>	0.25	0.25	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.25
		<i>N. gonorrhoeae</i>	0.5	-	-	-	<i>M. catarrhalis</i>	4	4	<i>N. gonorrhoeae</i>	ATCC® 49226	0.016-0.06
		<i>S. pneumoniae</i> (meningitis)	0.5	1	2	4	PK/PD (Non-species related) breakpoints	4	8	<i>H. influenzae</i>	ATCC® 49766	0.03-0.125
		<i>S. pneumoniae</i> (nonmeningitis)	1	2	4	-	-	-	-	-	-	-
		<i>Streptococcus</i> spp. β-Hemolytic Group	0.5	-	-	-	-	-	-	-	-	-
		<i>Streptococcus</i> spp. Viridans Group	1	2	4	-	-	-	-	-	-	-
		-	-	-	-	-	-	-	-	-	-	-
		CEFIXIME 0.016 - 256 µg/mL	CFM	Enterobacteriaceae	1	-	2	4	Enterobacteriaceae	1	1	<i>S. aureus</i>
<i>Haemophilus</i> spp.	1			-	-	-	<i>H. influenzae</i>	0.12	0.12	<i>E. coli</i>	ATCC® 25922	0.25-1
<i>N. gonorrhoeae</i>	0.25			-	-	-	<i>M. catarrhalis</i>	0.5	1	<i>H. influenzae</i>	ATCC® 49247	0.12-1
-	-	-	-	-	-	-	-	-	<i>N. gonorrhoeae</i>	ATCC® 49226	0.004-0.03	
-	-	-	-	-	-	-	-	-	<i>H. influenzae</i>	ATCC® 49766	0.016-0.06	
CEFOTAXIME 0.016 - 256 µg/mL 0.002 - 32 µg/mL	CTX	Enterobacteriaceae	1	-	2	4	Enterobacteriaceae	1	2	<i>S. aureus</i>	ATCC® 29213	1-4
		<i>Acinetobacter</i> spp.	8	-	16-32	64	<i>S. pneumoniae</i>	0.5	2	<i>E. coli</i>	ATCC® 25922	0.03-0.12
		Other Non-Enterobacteriaceae	8	-	16-32	64	Viridans group streptococci	0.5	0.5	<i>P. aeruginosa</i>	ATCC® 27853	8-32
		<i>Staphylococcus</i> spp.	8	-	16-32	64	<i>H. influenzae</i>	0.12	0.12	<i>H. influenzae</i>	ATCC® 49247	0.12-0.5
		<i>Haemophilus</i> spp.	2	-	-	-	<i>M. catarrhalis</i>	1	2	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.12
		<i>N. gonorrhoeae</i>	0.5	-	-	-	<i>N. gonorrhoeae</i>	0.12	0.12	<i>N. gonorrhoeae</i>	ATCC® 49226	0.016-0.06
		<i>S. pneumoniae</i> (meningitis)	0.5	1	2	4	<i>N. meningitidis</i>	0.12	0.12	<i>H. influenzae</i>	ATCC® 49766	0.004-0.016
		<i>S. pneumoniae</i> (nonmeningitis)	1	2	4	-	-	-	-	-	-	-
		<i>Streptococcus</i> spp. β-Hemolytic Group	0.5	-	-	-	<i>P. multocida</i>	0.03	0.03	-	-	-
		<i>Streptococcus</i> spp. Viridans Group	1	2	4	-	-	-	-	-	-	-
		<i>N. meningitidis</i>	0.12	-	-	-	<i>Kingella kingae</i>	0.12	0.12	-	-	-
		<i>Anaerobes</i>	16	32	64	-	-	-	-	-	-	-
CEFOTETAN 0.016 - 256 µg/mL	CTT	Enterobacteriaceae	16	-	32	64	Not available	-	-	<i>S. aureus</i>	ATCC® 29213	4-16
		<i>Staphylococcus</i> spp.	16	-	32	64	-	-	<i>E. coli</i>	ATCC® 25922	0.06-0.25	
		<i>N. gonorrhoeae</i>	2	-	4	8	-	-	<i>N. gonorrhoeae</i>	ATCC® 49226	0.5-2	
		Anaerobes	16	-	32	64	-	-	<i>B. fragilis</i>	ATCC® 25285	1-8	
-	-	-	-	-	-	-	-	<i>B. thetaiotaomicron</i>	ATCC® 29741	16-128		
-	-	-	-	-	-	-	-	<i>E. lenta</i>	ATCC® 43055	16-64		
CEFOXITIN 0.016 - 256 µg/mL	FOX	Enterobacteriaceae	8	-	16	32	<i>S. aureus</i> and <i>S. lugdunensis</i>	4	4	<i>S. aureus</i>	ATCC® 29213	1-4
		<i>Staphylococcus</i> spp.	4	-	-	8	(Methicillin/Oxacillin Resistant Staphylococci)	-	-	<i>E. coli</i>	ATCC® 25922	2-8
		<i>N. gonorrhoeae</i>	2	-	4	8	-	-	<i>N. gonorrhoeae</i>	ATCC® 49226	0.5-2	
		Anaerobes	16	-	32	64	-	-	<i>B. fragilis</i>	ATCC® 25285	2-8	
-	-	-	-	-	-	-	-	<i>B. thetaiotaomicron</i>	ATCC® 29741	8-64		
-	-	-	-	-	-	-	-	<i>E. lenta</i>	ATCC® 43055	2-16		
CEFPIROME 0.016 - 256 µg/mL	CR	Not available	-	-	-	-	Not available	-	-	<i>H. influenzae</i>	ATCC® 49247	0.25-1
CEFPODOXIME 0.016 - 256 µg/mL	PX	Enterobacteriaceae	2	-	4	8	Enterobacteriaceae	1	1	<i>S. aureus</i>	ATCC® 29213	1-8
		<i>Staphylococcus</i> spp.	2	-	4	8	<i>S. pneumoniae</i>	0.25	0.5	<i>E. coli</i>	ATCC® 25922	0.25-1
		<i>Haemophilus</i> spp.	2	-	-	-	<i>H. influenzae</i>	0.25	0.5	<i>H. influenzae</i>	ATCC® 49247	0.25-1
		<i>N. gonorrhoeae</i>	0.5	-	-	-	-	-	-	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.12
		<i>S. pneumoniae</i>	0.5	1	2	-	-	-	-	<i>N. gonorrhoeae</i>	ATCC® 49226	0.03-0.12
-	-	-	-	-	-	-	-	<i>H. influenzae</i>	ATCC® 49766	0.03-0.125		

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CEFTAROLINE 0.016 - 256 µg/mL 0.002 - 32 µg/mL	CPT	Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Haemophilus</i> spp. <i>S. pneumoniae</i> (nonmeningitis) <i>Streptococcus</i> spp. β-Hemolytic Group	0.5 1 0.5 0.5 0.5		1 2 - - -	2 4 - - -	Enterobacteriaceae <i>S. aureus</i> <i>S. pneumoniae</i> <i>H. influenzae</i> PK/PD (Non-species related) breakpoints	0.5 1 0.25 0.03 0.5	0.5 1 0.25 0.03 0.5	<i>S. aureus</i> ATCC® 29213 <i>E. faecalis</i> ATCC® 29212 <i>E. coli</i> ATCC® 25922 <i>K. pneumoniae</i> ATCC® 700603 <i>H. influenzae</i> ATCC® 49247 <i>S. pneumoniae</i> ATCC® 49619 <i>B. fragilis</i> ATCC® 25285 <i>B. thetaiotaomicron</i> ATCC® 29741 <i>C. difficile</i> ATCC® 700057 <i>H. influenzae</i> ATCC® 49766	0.12-0.5 0.25-2 0.03-0.12 2-8 0.03-0.12 0.008-0.03 2-16 8-64 0.5-4 0.004-0.016
CEFTAZIDIME 0.016 - 256 µg/mL	CAZ	Enterobacteriaceae <i>P. aeruginosa</i> <i>Acinetobacter</i> spp. <i>B. cepacia</i> <i>S. maltophilia</i> Other Non-Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Haemophilus</i> spp. <i>N. gonorrhoeae</i>	4 8 8 8 8 8 2 0.5		8 16 16 16 16 16 - -	16 32 32 32 32 32 - -	Enterobacteriaceae <i>Pseudomonas</i> spp. PK/PD (Non-species related) breakpoints	1 8 4	4 8 8	<i>S. aureus</i> ATCC® 29213 <i>E. coli</i> ATCC® 25922 <i>P. aeruginosa</i> ATCC® 27853 <i>K. pneumoniae</i> ATCC® 700603 <i>H. influenzae</i> ATCC® 49247 <i>N. gonorrhoeae</i> ATCC® 49226	4-16 0.06-0.5 1-4 16-64 0.12-1 0.03-0.12
CEFTAZIDIME*- AVIBACTAM (4 µg/mL) 0.016 - 256 * µg/mL	CZA	Not available					Enterobacteriaceae <i>Pseudomonas</i> spp. PK/PD (Non-species related) breakpoints	8 8 8	8 8 8	<i>S. aureus</i> ATCC® 29213 <i>E. coli</i> ATCC® 25922 <i>P. aeruginosa</i> ATCC® 27853 <i>E. coli</i> ATCC® 35218 <i>K. pneumoniae</i> ATCC® 700603 <i>H. influenzae</i> ATCC® 49247 <i>H. influenzae</i> ATCC® 49766 <i>S. pneumoniae</i> ATCC® 49619	4-16 0.06-0.5 0.5-4 0.03-0.12 0.25-2 0.06-0.5 0.016-0.06 0.25-2
CEFTIBUTEN 0.002 - 32 µg/mL	CTB	Enterobacteriaceae <i>H. influenzae</i>	8 2		16 -	32 -	Enterobacteriaceae (UTI only) <i>H. influenzae</i>	1 1	1 1	<i>E. coli</i> ATCC® 25922 <i>H. influenzae</i> ATCC® 49247 <i>H. influenzae</i> ATCC® 49766	0.12-0.5 0.25-1 0.016-0.06
CEFTIZOXIME 0.002 - 32 µg/mL 0.016 - 256 µg/mL	CZX	Enterobacteriaceae Other Non-Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Haemophilus</i> spp. <i>N. gonorrhoeae</i> <i>Anaerobes</i>	1 8 8 2 0.5 32		2 16-32 16-32 - - 64	4 64 64 - - 128	Not available			<i>S. aureus</i> ATCC® 29213 <i>E. coli</i> ATCC® 25922 <i>P. aeruginosa</i> ATCC® 27853 <i>H. influenzae</i> ATCC® 49247 <i>S. pneumoniae</i> ATCC® 49619 <i>N. gonorrhoeae</i> ATCC® 49226 <i>E. lenta</i> ATCC® 43055	2-8 0.03-0.12 16-64 0.06-0.5 0.12-0.5 0.008-0.03 8-32
CEFTOBIPROLE 0.002 - 32 µg/mL	BPR	Not available					Enterobacteriaceae <i>S. aureus</i> <i>S. pneumoniae</i> PK/PD (Non-species related) breakpoints	0.25 2 0.5 4	0.25 2 0.5 4	<i>S. aureus</i> ATCC® 29213 <i>E. faecalis</i> ATCC® 29212 <i>E. coli</i> ATCC® 25922 <i>P. aeruginosa</i> ATCC® 27853 <i>H. influenzae</i> ATCC® 49247 <i>H. influenzae</i> ATCC® 49766 <i>S. pneumoniae</i> ATCC® 49619	0.12-1 0.06-0.5 0.03-0.12 1-4 0.12-1 0.016-0.06 0.004-0.03
CEFTOLOZANE*- TAZOBACTAM (4 µg/mL) 0.016 - 256 * µg/mL	C/T	Enterobacteriaceae <i>P. aeruginosa</i> <i>Streptococcus</i> spp. Viridans Group	2 4 8		4 8 16	8 16 32	Enterobacteriaceae <i>P. aeruginosa</i>	1 4	1 4	<i>S. aureus</i> ATCC® 29213 <i>E. coli</i> ATCC® 25922 <i>P. aeruginosa</i> ATCC® 27853 <i>E. coli</i> ATCC® 35218 <i>K. pneumoniae</i> ATCC® 700603	16-64 0.12-0.5 0.25-1 0.06-0.25 0.5-2

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CEFTRIAXONE 0.016 - 256 µg/mL 0.002 - 32 µg/mL	CRO	Enterobacteriaceae	1		2	4	Enterobacteriaceae	1	2	<i>S. aureus</i>	ATCC® 29213	1-8
		<i>Acinetobacter spp.</i>	8		16-32	64	<i>S. pneumoniae</i>	0.5	2	<i>E. coli</i>	ATCC® 25922	0.03-0.12
		Other Non-Enterobacteriaceae	8		16-32	64	Viridans group streptococci	0.5	0.5	<i>P. aeruginosa</i>	ATCC® 27853	8-64
		<i>Staphylococcus spp.</i>	8		16-32	64	<i>H. influenzae</i>	0.12	0.12	<i>H. influenzae</i>	ATCC® 49247	0.06-0.25
		<i>Haemophilus spp.</i>	2		-	-	<i>M. catarrhalis</i>	1	2	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.12
		<i>N. gonorrhoeae</i>	0.25		-	-	<i>N. gonorrhoeae</i>	0.12	0.12	<i>N. gonorrhoeae</i>	ATCC® 49226	0.004-0.016
		<i>S. pneumoniae</i> (meningitis)	0.5		1	2	<i>N. meningitidis</i>	0.12	0.12	<i>H. influenzae</i>	ATCC® 49766	0.002-0.008
		<i>S. pneumoniae</i> (nonmeningitis)	1		2	4	<i>Kingella kingae</i>	0.06	0.06			
		<i>Streptococcus spp. β-Hemolytic Group</i>	0.5		-	-	PK/PD (Non-species related) breakpoints	1	2			
		<i>Streptococcus spp. Viridans Group</i>	1			2	4					
		<i>N. meningitidis</i>	0.12		-	-						
		Anaerobes	16			32	64					
		CEFUROXIME 0.016 - 256 µg/mL	CXM	Enterobacteriaceae (parenteral)	8		16	32	Enterobacteriaceae	8	8	<i>S. aureus</i>
Enterobacteriaceae (oral)	4				8-16	32	<i>S. pneumoniae</i> (iv)	0.5	1	<i>E. coli</i>	ATCC® 25922	2-8
<i>Haemophilus spp</i>	4				8	16	<i>S. pneumoniae</i> (oral)	0.25	0.5	<i>H. influenzae</i>	ATCC® 49766	0.25-1
<i>N. gonorrhoeae</i>	1				2	4	Viridans group streptococci (iv)	0.5	0.5	<i>S. pneumoniae</i>	ATCC® 49619	0.25-1
<i>S. pneumoniae</i> (parenteral)	0.5				1	2	<i>H. influenzae</i> (iv)	1	2	<i>N. gonorrhoeae</i>	ATCC® 49226	0.25-1
<i>S. pneumoniae</i> (oral)	1				2	4	<i>H. influenzae</i> (oral)	0.12	1			
							<i>M. catarrhalis</i> (iv)	4	8			
					<i>M. catarrhalis</i> (oral)	0.12	4					
					<i>Kingella kingae</i> (iv)	0.5	0.5					
					PK/PD (Non-species related) breakpoints (iv)	4	8					
CEPHALOTHIN 0.016 - 256 µg/mL	KF	Not available					Not available			<i>S. aureus</i>	ATCC® 29213	0.12-0.5
										<i>E. coli</i>	ATCC® 25922	4-16
										<i>S. pneumoniae</i>	ATCC® 49619	0.5-2
CHLORAMPHENICOL 0.016 - 256 µg/mL	C	Enterobacteriaceae	8		16	32	Enterobacteriaceae	8	8	<i>S. aureus</i>	ATCC® 29213	2-16
		<i>B. cepacia</i>	8		16	32	<i>Staphylococcus spp.</i>	8	8	<i>E. faecalis</i>	ATCC® 29212	4-16
		<i>S. maltophilia</i>	8		16	32	Streptococcus groups A, B, C and G	8	8	<i>E. coli</i>	ATCC® 25922	2-8
		Other non Enterobacteriaceae	8		16	32	<i>S. pneumoniae</i>	8	8	<i>H. influenzae</i>	ATCC® 49247	0.25-1
		<i>Staphylococcus spp.</i>	8		16	32	<i>H. influenzae</i>	2	2	<i>S. pneumoniae</i>	ATCC® 49619	2-8
		<i>Enterococcus spp.</i>	8		16	32	<i>M. catarrhalis</i>	2	2	<i>B. fragilis</i>	ATCC® 25285	4-16
		<i>Haemophilus spp.</i>	2		4	8	<i>N. meningitidis</i>	2	4	<i>B. thetaiotaomicron</i>	ATCC® 29741	8-32
		<i>S. pneumoniae</i>	4		-	8	Gram-positive anaerobes except <i>C. difficile</i>	8	8	<i>E. lenta</i>	ATCC® 43055	4-16
		<i>Streptococcus spp. β-Hemolytic Group</i>	4		8	16	Gram-negative anaerobes	8	8	<i>H. influenzae</i>	ATCC® 49766	0.25-1
		<i>Streptococcus spp. Viridans Group</i>	4			8	16					
		<i>N. meningitidis</i>	2		4	8						
		Anaerobes	8		16	32						

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
CIPROFLOXACIN 0.002 - 32 µg/mL	CIP	Enterobacteriaceae except	1		2	4	Enterobacteriaceae	0.25	0.5	<i>S. aureus</i>	ATCC® 29213	0.12-0.5
		<i>Salmonella</i> spp.					<i>Salmonella</i> spp.	0.06	0.06	<i>E. faecalis</i>	ATCC® 29212	0.25-2
		<i>Salmonella</i> spp.	0.06		0.12-0.5	1	<i>Pseudomonas</i> spp.	0.5	0.5	<i>E. coli</i>	ATCC® 25922	0.004-0.016
		<i>P. aeruginosa</i>	1		2	4	<i>Acinetobacter</i> spp.	1	1	<i>P. aeruginosa</i>	ATCC® 27853	0.25-1
		<i>Acinetobacter</i> spp.	1		2	4	<i>Staphylococcus</i> spp.	1	1	<i>H. influenzae</i>	ATCC® 49247	0.004-0.03
		Other non Enterobacteriaceae	1		2	4	<i>Enterococcus</i> spp.	4	4	<i>N. gonorrhoeae</i>	ATCC® 49226	0.001-0.008
		<i>Staphylococcus</i> spp.	1		2	4	<i>S. pneumoniae</i>	0.12	2	<i>H. influenzae</i>	ATCC® 49766	0.004-0.016
		<i>Enterococcus</i> spp.	1		2	4	<i>H. influenzae</i>	0.06	0.06			
		<i>Haemophilus</i> spp.	1		-	-	<i>M. catarrhalis</i>	0.5	0.5			
		<i>N. gonorrhoeae</i>	0.06		0.12-0.5	1	<i>N. gonorrhoeae</i>	0.03	0.06			
		<i>N. meningitidis</i>	0.03		0.06	0.12	<i>N. meningitidis</i>	0.03	0.03			
							<i>P. multocida</i>	0.06	0.06			
							<i>C. jejuni</i> and <i>coli</i>	0.5	0.5			
							<i>Corynebacterium</i> spp.	1	1			
					<i>Aerococcus sanguinicola</i> and <i>urinae</i>	2	2					
					<i>Kingella kingae</i>	0.06	0.06					
					PK/PD (Non-species related) breakpoints	0.25	0.5					
CLARITHROMYCIN 0.016 - 256 µg/mL	CLR	<i>Staphylococcus</i> spp.	2		4	8	<i>Staphylococcus</i> spp.	1	2	<i>S. aureus</i>	ATCC® 29213	0.12-0.5
		<i>H. influenzae</i>	8		16	32	Streptococcus groups A, B, C and G	0.25	0.5	<i>H. influenzae</i> (-CO <sub>2</sub> )	ATCC® 49247	4-16
		<i>S. pneumoniae</i>	0.25		0.5	1	<i>S. pneumoniae</i>	0.25	0.5	<i>H. influenzae</i> (+CO <sub>2</sub> )	ATCC® 49247	8-32
		Streptococcus spp. β-Hemolytic Group	0.25		0.5	1	<i>H. influenzae</i>	1	32	<i>S. pneumoniae</i> (-CO <sub>2</sub> )	ATCC® 49619	0.032-0.125
		Streptococcus spp. Viridans Group	0.25		0.5	1	<i>M. catarrhalis</i>	0.25	0.5	<i>S. pneumoniae</i> (+CO <sub>2</sub> )	ATCC® 49619	0.064-0.25
							<i>H. pylori</i>	0.25	0.5			
							<i>Kingella kingae</i>	0.5	0.5			
CLINDAMYCIN 0.016 - 256 µg/mL	CD	<i>Staphylococcus</i> spp.	0.5		1-2	4	<i>Staphylococcus</i> spp.	0.25	0.5	<i>S. aureus</i>	ATCC® 29213	0.06-0.25
		<i>S. pneumoniae</i>	0.25		0.5	1	Streptococcus groups A, B, C and G	0.5	0.5	<i>E. faecalis</i>	ATCC® 29212	4-16
		Streptococcus spp. β-Hemolytic Group	0.25		0.5	1	<i>S. pneumoniae</i>	0.5	0.5	<i>S. pneumoniae</i> (-CO <sub>2</sub> )	ATCC® 49619	0.032-0.125
		Streptococcus spp. Viridans Group	0.25		0.5	1	Viridans group streptococci	0.5	0.5	<i>S. pneumoniae</i> (+CO <sub>2</sub> )	ATCC® 49619	0.064-0.25
		Anaerobes	2		4	8	Gram-positive anaerobes except <i>C. difficile</i>	4	4	<i>B. fragilis</i>	ATCC® 25285	0.5-2
							Gram-negative anaerobes	4	4	<i>B. thetaiotaomicron</i>	ATCC® 29741	2-8
							<i>Corynebacterium</i> spp.	0.5	0.5	<i>E. lenta</i>	ATCC® 43055	0.06-0.25
COLISTIN 0.016 - 256 µg/mL 0.064 - 1024 µg/mL	CS	<i>P. aeruginosa</i>	2		-	4	Enterobacteriaceae	2	2	<i>E. coli</i>	ATCC® 25922	0.25-2
		<i>Acinetobacter</i> spp.	2		-	4	<i>Pseudomonas</i> spp.	2	2	<i>P. aeruginosa</i>	ATCC® 27853	0.5-4
DALBAVANCIN 0.002 - 32 µg/mL	DAL	Not available					<i>Staphylococcus</i> spp.	0.12	0.12	<i>S. aureus</i>	ATCC® 29213	0.03-0.12
										<i>E. faecalis</i>	ATCC® 29212	0.03-0.12
										<i>S. pneumoniae</i>	ATCC® 49619	0.008-0.03
DAPTOMYCIN 0.016 - 256 µg/mL	DAP	<i>Staphylococcus</i> spp.	1		-	-	<i>Staphylococcus</i> spp.	1	1	<i>S. aureus</i>	ATCC® 29213	0.12-1
		<i>Enterococcus</i> spp.	4		-	-	Streptococcus groups A, B, C and G	1	1	<i>E. faecalis</i>	ATCC® 29212	1-4
		Streptococcus spp. β-Hemolytic Group	1		-	-				<i>S. pneumoniae</i>	ATCC® 49619	0.06-0.5
		Streptococcus spp. Viridans Group	1		-	-						

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
DORIPENEM 0.002 - 32 µg/mL	DOR	Enterobacteriaceae	1		2	4	Enterobacteriaceae	1	2	<i>S. aureus</i>	ATCC® 29213	0.016-0.06
		<i>P. aeruginosa</i>	2		4	8	<i>Pseudomonas</i> spp.	1	2	<i>E. faecalis</i>	ATCC® 29212	1-4
		<i>Acinetobacter</i> spp.	2		4	8	<i>Acinetobacter</i> spp.	1	2	<i>E. coli</i>	ATCC® 25922	0.016-0.06
		<i>Staphylococcus</i> spp.	0.5		-	-	<i>S. pneumoniae</i>	1	1	<i>P. aeruginosa</i>	ATCC® 27853	0.12-0.5
		<i>Haemophilus</i> spp.	1		-	-	Viridans group streptococci	1	1	<i>H. influenzae</i>	ATCC® 49766	0.06-0.25
		<i>S. pneumoniae</i>	1		-	-	<i>H. influenzae</i>	1	1	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.12
		<i>Streptococcus</i> spp. β-Hemolytic Group	0.12		-	-	<i>M. catarrhalis</i>	1	1	<i>B. fragilis</i>	ATCC® 25285	0.12-0.5
		<i>Streptococcus</i> spp. Viridans Group	1		-	-	Gram-positive anaerobes except <i>C. difficile</i>	1	1	<i>B. thetaiotaomicron</i>	ATCC® 29741	0.12-1
		Anaerobes	2		4	8	Gram-negative anaerobes	1	1			
							PK/PD (Non-species related) breakpoints	1	2			
DOXYCYCLINE 0.016 - 256 µg/mL	DXT	Enterobacteriaceae	4		8	16	<i>Staphylococcus</i> spp.	1	2	<i>S. aureus</i>	ATCC® 29213	0.12-0.5
		<i>Acinetobacter</i> spp.	4		8	16	Streptococcus groups A, B, C and G	1	2	<i>E. faecalis</i>	ATCC® 29212	2-8
		Other Non-Enterobacteriaceae	4		8	16	<i>S. pneumoniae</i>	1	2	<i>E. coli</i>	ATCC® 25922	0.5-2
		<i>Staphylococcus</i> spp.	4		8	16	<i>H. influenzae</i>	1	2	<i>S. pneumoniae</i>	ATCC® 49619	0.016-0.12
		<i>Enterococcus</i> spp.	4		8	16	<i>M. catarrhalis</i>	1	2	<i>B. thetaiotaomicron</i>	ATCC® 29741	2-8
		<i>S. pneumoniae</i>	0.25		0.5	1	<i>P. multocida</i>	1	1	<i>E. lenta</i>	ATCC® 43055	2-16
							<i>Kingella kingae</i>	0.5	0.5	<i>H. influenzae</i>	ATCC® 49766	0.25-1
ENROFLOXACIN 0.002 - 32 µg/mL	ENR	Refer to CLSI VET01S					Not available			<i>S. aureus</i>	ATCC® 29213	0.03-0.12
										<i>E. faecalis</i>	ATCC® 29212	0.12-1
										<i>E. coli</i>	ATCC® 25922	0.008-0.03
										<i>P. aeruginosa</i>	ATCC® 27853	1-4
ERTAPENEM 0.002 - 32 µg/mL	ETP	Enterobacteriaceae	0.5		1	2	Enterobacteriaceae	0.5	1	<i>S. aureus</i>	ATCC® 29213	0.06-0.25
		<i>Staphylococcus</i> spp.	2		4	8	<i>S. pneumoniae</i>	0.5	0.5	<i>E. faecalis</i>	ATCC® 29212	4-16
		<i>Haemophilus</i> spp.	0.5		-	-	Viridans group streptococci	0.5	0.5	<i>E. coli</i>	ATCC® 25922	0.004-0.016
		<i>S. pneumoniae</i>	1		2	4	<i>H. influenzae</i>	0.5	0.5	<i>P. aeruginosa</i>	ATCC® 27853	2-8
		<i>Streptococcus</i> spp. β-Hemolytic Group	1		-	-	<i>M. catarrhalis</i>	0.5	0.5	<i>H. influenzae</i>	ATCC® 49766	0.016-0.06
		<i>Streptococcus</i> spp. Viridans Group	1		-	-	Gram-positive anaerobes except <i>C. difficile</i>	1	1	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.25
		Anaerobes	4		8	16	Gram-negative anaerobes	1	1	<i>B. fragilis</i>	ATCC® 25285	0.06-0.5
							PK/PD (Non-species related) breakpoints	0.5	1	<i>B. thetaiotaomicron</i>	ATCC® 29741	0.5-2
								<i>E. lenta</i>	ATCC® 43055	0.5-4		
ERYTHROMYCIN 0.016 - 256 µg/mL	E	<i>Staphylococcus</i> spp.	0.5		1-4	8	<i>Staphylococcus</i> spp.	1	2	<i>S. aureus</i>	ATCC® 29213	0.25-1
		<i>Enterococcus</i> spp.	0.5		1-4	8	Streptococcus groups A, B, C and G	0.25	0.5	<i>E. faecalis</i>	ATCC® 29212	1-4
		<i>S. pneumoniae</i>	0.25		0.5	1	<i>S. pneumoniae</i>	0.25	0.5	<i>S. pneumoniae</i> (-CO <sub>2</sub> )	ATCC® 49619	0.032-0.125
		<i>Streptococcus</i> spp. β-Hemolytic Group	0.25		0.5	1	<i>H. influenzae</i>	0.5	16	<i>S. pneumoniae</i> (+CO <sub>2</sub> )	ATCC® 49619	0.064-0.25
		<i>Streptococcus</i> spp. Viridans Group	0.25		0.5	1	<i>M. catarrhalis</i>	0.25	0.5			
							<i>L. monocytogenes</i>	1	1			
FLORFENICOL 0.016 - 256 µg/mL	FFC	Refer to CLSI VET01S					Not available			<i>S. aureus</i>	ATCC® 29213	2-8
										<i>E. faecalis</i>	ATCC® 29212	2-8
										<i>E. coli</i>	ATCC® 25922	2-8
										<i>S. pneumoniae</i>	ATCC® 49619	1-4
FOSFOMYCIN 0.016 - 256 µg/mL 0.064 - 1024 µg/mL	FOS	Enterobacteriaceae	64		128	256	Enterobacteriaceae	32	32	<i>S. aureus</i>	ATCC® 29213	0.5-4
		<i>Enterococcus</i> spp.	64		128	256	<i>Staphylococcus</i> spp (iv)	32	32	<i>E. faecalis</i>	ATCC® 29212	32-128
										<i>E. coli</i>	ATCC® 25922	0.5-2
										<i>P. aeruginosa</i>	ATCC® 27853	2-8
FUSIDIC ACID 0.016 - 256 µg/mL	FU	Not available					<i>Staphylococcus</i> spp.			<i>S. aureus</i>	ATCC® 29213	0.06-0.25
										<i>S. pneumoniae</i>	ATCC® 49619	4-32

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
GATIFLOXACIN 0.002 - 32 µg/mL	GAT	Enterobacteriaceae	2		4	8	Not available			<i>S. aureus</i>	ATCC® 29213	0.03-0.12
		<i>P. aeruginosa</i>	2		4	8		<i>E. faecalis</i>	ATCC® 29212	0.12-1		
		<i>Acinetobacter</i> spp.	2		4	8		<i>E. coli</i>	ATCC® 25922	0.008-0.03		
		Other Non-Enterobacteriaceae	2		4	8		<i>P. aeruginosa</i>	ATCC® 27853	0.5-2		
		<i>Staphylococcus</i> spp.	0.5		1	2		<i>H. influenzae</i>	ATCC® 49766	0.004-0.03		
		<i>Enterococcus</i> spp.	2		4	8		<i>S. pneumoniae</i>	ATCC® 49619	0.12-0.5		
		<i>Haemophilus</i> spp.	1		-	-		<i>N. gonorrhoeae</i>	ATCC® 49226	0.002-0.016		
		<i>N. gonorrhoeae</i>	0.125		0.25	0.5						
		<i>S. pneumoniae</i>	1		2	4						
		<i>Streptococcus</i> spp. β-Hemolytic Group	1		2	4						
		<i>Streptococcus</i> spp. Viridans Group	1		2	4						
GEMIFLOXACIN 0.002 - 32 µg/mL	GEM	Enterobacteriaceae	0.25		0.5	1	Not available			<i>S. aureus</i>	ATCC® 29213	0.008-0.03
		<i>Haemophilus</i> spp.	0.12		-	-		<i>E. faecalis</i>	ATCC® 29212	0.016-0.12		
		<i>S. pneumoniae</i>	0.12		0.25	0.5		<i>E. coli</i>	ATCC® 25922	0.004-0.016		
								<i>P. aeruginosa</i>	ATCC® 27853	0.25-1		
									<i>H. influenzae</i>	ATCC® 49766	0.002-0.008	
									<i>S. pneumoniae</i>	ATCC® 49619	0.008-0.03	
GENTAMICIN 0.016 - 256 µg/mL 0.064 - 1024 µg/mL	CN	Enterobacteriaceae	4		8	16	Enterobacteriaceae	2	4	<i>S. aureus</i>	ATCC® 29213	0.12-1
		<i>P. aeruginosa</i>	4		8	16	<i>Pseudomonas</i> spp.	4	4	<i>E. faecalis</i>	ATCC® 29212	4-16
		<i>Acinetobacter</i> spp.	4		8	16	<i>Acinetobacter</i> spp.	4	4	<i>E. coli</i>	ATCC® 25922	0.25-1
		Other Non-Enterobacteriaceae	4		8	16	<i>S. aureus</i>	1	1	<i>P. aeruginosa</i>	ATCC® 27853	0.5-2
		<i>Staphylococcus</i> spp.	4		8	16	Coagulase-negative Staphylococci	1	1	HLAR		
		<i>Enterococcus</i> spp. (screening test for HLAR)	500		-	500	<i>Corynebacterium</i> spp.	1	1	<i>E. faecalis</i>	ATCC® 29212	≤ 500
							PK/PD (Non-species related) breakpoints	2	4	<i>E. faecalis</i>	ATCC® 51299	> 500
							<i>Enterococcus</i> spp. (test for high-level gentamicin resistance)	128	128			
IMIPENEM 0.002 - 32 µg/mL	IMI	Enterobacteriaceae	1		2	4	Enterobacteriaceae	2	8	<i>S. aureus</i>	ATCC® 29213	0.016-0.06
		<i>P. aeruginosa</i>	2		4	8	<i>Pseudomonas</i> spp.	4	8	<i>E. faecalis</i>	ATCC® 29212	0.5-2
		<i>Acinetobacter</i> spp.	2		4	8	<i>Acinetobacter</i> spp.	2	8	<i>E. coli</i>	ATCC® 25922	0.06-0.25
		Other Non-Enterobacteriaceae	4		8	16	<i>Enterococcus</i> spp.	4	8	<i>P. aeruginosa</i>	ATCC® 27853	1-4
		<i>Staphylococcus</i> spp.	4		8	16	<i>S. pneumoniae</i>	2	2	<i>H. influenzae</i>	ATCC® 49766	0.25-1
		<i>Haemophilus</i> spp.	4		-	-	Viridans group streptococci	2	2	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.12
		<i>S. pneumoniae</i>	0.12		0.25-0.5	1	<i>H. influenzae</i>	2	2	<i>B. fragilis</i>	ATCC® 25285	0.03-0.25
		Anaerobes	4		8	16	<i>M. catarrhalis</i>	2	2	<i>B. thetaiotaomicron</i>	ATCC® 29741	0.25-1
							Gram-positive anaerobes except <i>C. difficile</i>	2	8	<i>E. lenta</i>	ATCC® 43055	0.25-2
							Gram-negative anaerobes	2	8			
							PK/PD (Non-species related) breakpoints	2	8			
KANAMYCIN 0.016 - 256 µg/mL	K	Enterobacteriaceae	16		32	64	Not available			<i>S. aureus</i>	ATCC® 29213	1-4
		<i>Staphylococcus</i> spp.	16		32	64		<i>E. faecalis</i>	ATCC® 29212	16-64		
								<i>E. coli</i>	ATCC® 25922	1-4		



ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
LEVOFLOXACIN 0.002 - 32 µg/mL	LEV	Enterobacteriaceae	0.12		0.25-1	2	Enterobacteriaceae	0.5	1	<i>S. aureus</i>	ATCC® 29213	0.06-0.5
		<i>P. aeruginosa</i>	2		4	8	<i>Pseudomonas</i> spp.	1	1	<i>E. faecalis</i>	ATCC® 29212	0.25-2
		<i>Acinetobacter</i> spp.	2		4	8	<i>Acinetobacter</i> spp.	0.5	1	<i>E. coli</i>	ATCC® 25922	0.008-0.06
		<i>B. cepacia</i>	2		4	8	<i>Staphylococcus</i> spp.	1	1	<i>P. aeruginosa</i>	ATCC® 27853	0.5-4
		<i>S. maltophilia</i>	2		4	8	<i>Enterococcus</i> spp.	4	4	<i>H. influenzae</i>	ATCC® 49247	0.008-0.03
		Other Non-Enterobacteriaceae	2		4	8	<i>Streptococcus</i> groups A, B, C and G	2	2	<i>S. pneumoniae</i>	ATCC® 49619	0.5-2
		<i>Staphylococcus</i> spp.	1		2	4	<i>S. pneumoniae</i>	2	2	<i>H. influenzae</i>	ATCC® 49766	0.008-0.03
		<i>Enterococcus</i> spp.	2		4	8	<i>H. influenzae</i>	0.06	0.06			
		<i>Haemophilus</i> spp.	2		-	-	<i>M. catarrhalis</i>	1	1			
		<i>S. pneumoniae</i>	2		4	8	<i>H. pylori</i>	1	1			
		<i>Streptococcus</i> spp. β-Hemolytic Group	2		4	8	<i>P. multocida</i>	0.06	0.06			
		<i>Streptococcus</i> spp. Viridans Group	2		4	8	<i>Aerococcus sanguinicola</i> and <i>urinae</i>	2	2			
		<i>N. meningitidis</i>	0.03		0.06	0.12	<i>Kingella kingae</i>	0.125	0.125			
							PK/PD (Non-species related) breakpoints	0.5	1			
		LINEZOLID 0.016 - 256 µg/mL	LNZ	<i>Staphylococcus</i> spp.	4		-	8	<i>Staphylococcus</i> spp.	4	4	<i>S. aureus</i>
<i>Enterococcus</i> spp.	2				4	8	<i>Enterococcus</i> spp.	4	4	<i>E. faecalis</i>	ATCC® 29212	1-4
<i>S. pneumoniae</i>	2				-	-	<i>Streptococcus</i> groups A, B, C and G	2	4	<i>S. pneumoniae</i>	ATCC® 49619	0.25-2
<i>Streptococcus</i> spp. β-Hemolytic Group	2				-	-	<i>S. pneumoniae</i>	2	4	<i>B. fragilis</i>	ATCC® 25285	2-8
<i>Streptococcus</i> spp. Viridans Group	2				-	-	<i>Corynebacterium</i> spp.	2	2	<i>B. thetaiotaomicron</i>	ATCC® 29741	2-8
							PK/PD (Non-species related) breakpoints	2	4	<i>E. lenta</i>	ATCC® 43055	0.5-2
MARBOFLOXACIN 0.002 - 32 µg/mL	MAR	Refer to CLSI VET01S				Not available				<i>S. aureus</i>	ATCC® 29213	0.12-0.5
										<i>E. faecalis</i>	ATCC® 29212	0.5-2
										<i>E. coli</i>	ATCC® 25922	0.008-0.03
										<i>P. aeruginosa</i>	ATCC® 27853	0.5-2
MECILLINAM 0.016 - 256 µg/mL	MEC	Enterobacteriaceae	8		16	32	Enterobacteriaceae	8	8	<i>E. coli</i>	ATCC® 25922	0.03-0.25
MEROPENEM 0.002 - 32 µg/mL 0.016 - 256 µg/mL	MRP	Enterobacteriaceae	1		2	4	Enterobacteriaceae	2	8	<i>S. aureus</i>	ATCC® 29213	0.03-0.12
		<i>P. aeruginosa</i>	2		4	8	<i>Pseudomonas</i> spp.	2	8	<i>E. faecalis</i>	ATCC® 29212	2-8
		<i>Acinetobacter</i> spp.	2		4	8	<i>Acinetobacter</i> spp.	2	8	<i>E. coli</i>	ATCC® 25922	0.008-0.06
		<i>B. cepacia</i>	4		8	16	<i>S. pneumoniae</i> (infections other than meningitis)	2	2	<i>P. aeruginosa</i>	ATCC® 27853	0.12-1
		Other Non-Enterobacteriaceae	4		8	16	<i>S. pneumoniae</i> (meningitis)	0.25	1	<i>H. influenzae</i>	ATCC® 49766	0.03-0.12
		<i>Staphylococcus</i> spp.	4		8	16	Viridans group streptococci	2	2	<i>S. pneumoniae</i>	ATCC® 49619	0.03-0.25
		<i>Haemophilus</i> spp.	0.5		-	-	<i>H. influenzae</i> (infections other than meningitis)	2	2	<i>B. fragilis</i>	ATCC® 25285	0.03-0.25
		<i>S. pneumoniae</i>	0.25		0.5	1	<i>H. influenzae</i> (meningitis)	0.25	1	<i>B. thetaiotaomicron</i>	ATCC® 29741	0.06-0.5
		<i>Streptococcus</i> spp. β-Hemolytic Group	0.5		-	-	<i>M. catarrhalis</i>	2	2	<i>E. lenta</i>	ATCC® 43055	0.125-1
		<i>Streptococcus</i> spp. Viridans Group	0.5		-	-	<i>N. meningitidis</i>	0.25	0.25			
		<i>N. meningitidis</i>	0.25		-	-	Gram-positive anaerobes except <i>C. difficile</i>	2	8			
		Anaerobes	4		8	16	Gram-negative anaerobes	2	8			
							<i>L. monocytogenes</i>	0.25	0.25			
							<i>Kingella kingae</i>	0.03	0.03			
							PK/PD (Non-species related) breakpoints	2	8			
		METRONIDAZOLE 0.016 - 256 µg/mL	MTZ	Anaerobes	8		16	32	Gram-positive anaerobes except <i>C. difficile</i>	4	4	<i>B. fragilis</i>
							<i>C. difficile</i>	2	2	<i>B. thetaiotaomicron</i>	ATCC® 29741	0.5-4
							Gram-negative anaerobes	4	4	<i>E. lenta</i>	ATCC® 43055	0.125-0.5
							<i>H. pylori</i>	8	8			

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
MINOCYCLINE 0.016 - 256 µg/mL	MN	Enterobacteriaceae	4		8	16	<i>Staphylococcus</i> spp.	0.5	1	<i>S. aureus</i>	ATCC® 29213	0.06-0.5
		<i>Acinetobacter</i> spp.	4		8	16	<i>Streptococcus</i> groups A, B, C and G	0.5	1	<i>E. faecalis</i>	ATCC® 29212	1-4
		<i>B. cepacia</i>	4		8	16	<i>S. pneumoniae</i>	0.5	1	<i>E. coli</i>	ATCC® 25922	0.25-1
		<i>S. maltophilia</i>	4		8	16	<i>H. influenzae</i>	1	2	<i>H. influenzae</i>	ATCC® 49766	0.125-0.5
		Other Non-Enterobacteriaceae	4		8	16	<i>M. catarrhalis</i>	1	2			
		<i>Staphylococcus</i> spp.	4		8	16	<i>N. gonorrhoeae</i>	0.5	1			
		<i>Enterococcus</i> spp.	4		8	16	<i>N. meningitidis</i>	1	2			
		<i>N. meningitidis</i>	2		-	-						
MOXIFLOXACIN 0.002 - 32 µg/mL	MXF	<i>Staphylococcus</i> spp.	0.5		1	2	Enterobacteriaceae	0.25	0.25	<i>S. aureus</i>	ATCC® 29213	0.016-0.12
		<i>Haemophilus</i> spp.	1		-	-	<i>Staphylococcus</i> spp.	0.25	0.25	<i>E. faecalis</i>	ATCC® 29212	0.06-0.5
		<i>S. pneumoniae</i>	1		2	4	<i>Streptococcus</i> groups A, B, C and G	0.5	0.5	<i>E. coli</i>	ATCC® 25922	0.008-0.06
		Anaerobes	2		4	8	<i>S. pneumoniae</i>	0.5	0.5	<i>P. aeruginosa</i>	ATCC® 27853	1-8
							<i>H. influenzae</i>	0.125	0.125	<i>H. influenzae</i>	ATCC® 49247	0.008-0.03
							<i>M. catarrhalis</i>	0.5	0.5	<i>S. pneumoniae</i>	ATCC® 49619	0.06-0.25
							<i>Corynebacterium</i> spp.	0.5	0.5	<i>N. gonorrhoeae</i>	ATCC® 49226	0.008-0.03
							PK/PD (Non-species related) breakpoints	0.25	0.25	<i>B. fragilis</i>	ATCC® 25285	0.12-0.5
										<i>B. thetaiotaomicron</i>	ATCC® 29741	1-8
										<i>E. lenta</i>	ATCC® 43055	0.12-0.5
								<i>H. influenzae</i>	ATCC® 49766	0.008-0.03		
MUPIROCI 0.064 - 1024 µg/mL	MUP	<i>S. aureus</i> (screening test for HLMR)	256		-	256	<i>Staphylococcus</i> spp.	1	256	<i>S. aureus</i> (mupA negative)	ATCC® 29213	0.06-0.5
										<i>E. faecalis</i> (mupA negative)	ATCC® 29212	16-128
										<i>S. aureus</i> (mupA positive)	ATCC® BAA 1708	> 256
NALIDIXIC ACID 0.016 - 256 µg/mL	NA	Enterobacteriaceae	16		-	32	Not available			<i>E. coli</i>	ATCC® 25922	1-4
NETILMICIN 0.016 - 256 µg/mL	NET	Enterobacteriaceae	8		16	32	Enterobacteriaceae	2	4	<i>S. aureus</i>	ATCC® 29213	≤ 0.25
		<i>P. aeruginosa</i>	8		16	32	<i>Pseudomonas</i> spp.	4	4	<i>E. faecalis</i>	ATCC® 29212	4-16
		<i>Acinetobacter</i> spp.	8		16	32	<i>Acinetobacter</i> spp.	4	4	<i>E. coli</i>	ATCC® 25922	≤ 0.5-1
		Other Non-Enterobacteriaceae	8		16	32	<i>S. aureus</i>	1	1	<i>P. aeruginosa</i>	ATCC® 27853	0.5-8
		<i>Staphylococcus</i> spp.	8		16	32	Coagulase-negative Staphylococci	1	1			
							PK/PD (Non-species related) breakpoints	2	4			
NITROFURANTOIN 0.032 - 512 µg/mL	F	Enterobacteriaceae	32		64	128	Enterobacteriaceae	64	64	<i>S. aureus</i>	ATCC® 29213	8-32
		<i>Staphylococcus</i> spp.	32		64	128	<i>Staphylococcus</i> spp.	64	64	<i>E. faecalis</i>	ATCC® 29212	4-16
		<i>Enterococcus</i> spp.	32		64	128	<i>Enterococcus</i> spp.	64	64	<i>E. coli</i>	ATCC® 25922	4-16
							<i>Streptococcus</i> groups A, B, C and G	64	64	<i>S. pneumoniae</i>	ATCC® 49619	4-16
					<i>Aerococcus sanguinicola</i> and <i>urinae</i>	16	16					
NORFLOXACIN 0.016 - 256 µg/mL	NOR	Enterobacteriaceae	4		8	16	Enterobacteriaceae	0.5	1	<i>S. aureus</i>	ATCC® 29213	0.5-2
		<i>P. aeruginosa</i>	4		8	16				<i>E. faecalis</i>	ATCC® 29212	2-8
		Other Non-Enterobacteriaceae	4		8	16				<i>E. coli</i>	ATCC® 25922	0.03-0.12
		<i>Staphylococcus</i> spp.	4		8	16				<i>P. aeruginosa</i>	ATCC® 27853	1-4
		<i>Enterococcus</i> spp.	4		8	16				<i>S. pneumoniae</i>	ATCC® 49619	2-8
OFLOXACIN 0.002 - 32 µg/mL	OFX	Enterobacteriaceae	0.12		0.25-1	2	Enterobacteriaceae	0.25	0.5	<i>S. aureus</i>	ATCC® 29213	0.12-1
		<i>P. aeruginosa</i>	2		4	8	<i>Staphylococcus</i> spp.	1	1	<i>E. faecalis</i>	ATCC® 29212	1-4
		Other Non-Enterobacteriaceae	2		4	8	<i>S. pneumoniae</i>	0.12	4	<i>E. coli</i>	ATCC® 25922	0.016-0.12
		<i>Staphylococcus</i> spp.	1		2	4	<i>H. influenzae</i>	0.06	0.06	<i>P. aeruginosa</i>	ATCC® 27853	1-8
		<i>Haemophilus</i> spp.	2		-	-	<i>M. catarrhalis</i>	0.5	0.5	<i>H. influenzae</i>	ATCC® 49247	0.016-0.06
		<i>N. gonorrhoeae</i>	0.25		0.5-1	2	<i>N. gonorrhoeae</i>	0.12	0.25	<i>S. pneumoniae</i>	ATCC® 49619	1-4
		<i>S. pneumoniae</i>	2		4	8	PK/PD (Non-species related) breakpoints	0.25	0.5	<i>N. gonorrhoeae</i>	ATCC® 49226	0.004-0.016
		<i>Streptococcus</i> spp. β-Hemolytic Group	2		4	8				<i>H. influenzae</i>	ATCC® 49766	0.016-0.06
		<i>Streptococcus</i> spp. Viridans Group	2		4	8						

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
OXACILLIN 0.016 - 256 µg/mL	OX	<i>S. aureus</i> and <i>S. lugdunensis</i> Coagulase-negative Staphylococci except <i>S. lugdunensis</i> and <i>S. pseudintermedius</i> <i>S. pseudintermedius</i>	2 0.25 0.25	-	-	4 0.5 0.5	<i>S. aureus</i> and <i>S. lugdunensis</i> Coagulase-negative Staphylococci (Methicillin/Oxacillin Resistant Staphylococci)	2 0.25	2 0.25	<i>S. aureus</i> <i>E. faecalis</i>	ATCC® 29213 ATCC® 29212	0.12-0.5 8-32
PENICILLIN G 0.002 - 32 µg/mL or 0.016 - 256 µg/mL	P	<i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>N. gonorrhoeae</i> <i>S. pneumoniae</i> parenteral (nonmeningitis) <i>S. pneumoniae</i> parenteral (meningitis) <i>S. pneumoniae</i> oral <i>Streptococcus</i> spp. β-Hemolytic Group <i>Streptococcus</i> spp. Viridans Group <i>N. meningitidis</i> Anaerobes	0.12 8 0.06 2 0.06 0.12 0.12 0.06 0.12 0.5	- - 0.12-1 4 - - 0.12-1 -	- 16 2 8 0.12 - 2 - 1	0.25 0.5 2 8 0.12 - 2 - 2	<i>S. aureus</i> <i>Streptococcus</i> groups A, B, C and G <i>S. pneumoniae</i> (infections other than meningitis) <i>S. pneumoniae</i> (meningitis) Viridans group streptococci <i>N. gonorrhoeae</i> <i>N. meningitidis</i> Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes <i>Corynebacterium</i> spp. <i>Aerococcus sanguincola</i> and <i>urinae</i> <i>Kingella kingae</i> <i>L. monocytogenes</i> <i>P. multocida</i> PK/PD (Non-species related) breakpoints	0.12 0.25 0.06 0.25 0.06 0.06 0.25 0.12 0.12 0.03 1 0.5 0.25	0.12 0.25 2 0.06 2 0.25 0.5 0.12 0.12 0.03 1 0.5 2	<i>S. aureus</i> <i>E. faecalis</i> <i>S. pneumoniae</i> <i>N. gonorrhoeae</i> <i>B. fragilis</i> <i>B. thetaiotaomicron</i>	ATCC® 29213 ATCC® 29212 ATCC® 49619 ATCC® 49226 ATCC® 25285 ATCC® 29741	0.25-2 1-4 0.25-1 0.25-1 8-32 8-32
PIPERACILLIN 0.016 - 256 µg/mL	PIP	Enterobacteriaceae <i>P. aeruginosa</i> <i>Acinetobacter</i> spp. Other Non-Enterobacteriaceae Anaerobes	16 16 16 16 32	32-64	128	128	Enterobacteriaceae <i>Pseudomonas</i> spp. Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes PK/PD (Non-species related) breakpoints	8 16 8 16 4	16 16 16 16 16	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>E. coli</i> <i>B. fragilis</i> <i>B. thetaiotaomicron</i> <i>E. lenta</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 35218 ATCC® 25285 ATCC® 29741 ATCC® 43055	1-4 1-4 1-4 1-8 > 64 4-16 8-64 8-32
PIPERACILLIN* TAZOBACTAM (4 µg/mL) 0.016 - 256 * µg/mL	TZP	Enterobacteriaceae <i>P. aeruginosa</i> <i>Acinetobacter</i> spp. Other Non-Enterobacteriaceae <i>Haemophilus</i> spp. Anaerobes	16 16 16 16 1 32	32-64	128	128	Enterobacteriaceae <i>Pseudomonas</i> spp. Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes PK/PD (Non-species related) breakpoints	8 16 8 8 4	16 16 16 16 16	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>E. coli</i> <i>K. pneumoniae</i> <i>H. influenzae</i> <i>B. fragilis</i> <i>B. thetaiotaomicron</i> <i>E. lenta</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 35218 ATCC® 700603 ATCC® 49247 ATCC® 25285 ATCC® 29741 ATCC® 43055	0.25-2 1-4 1-4 1-8 0.5-2 8-32 0.06-0.5 0.03-0.25 2-16 8-32
POLYMYXIN B 0.064 - 1024 µg/mL	PB	<i>P. aeruginosa</i> <i>Acinetobacter</i> spp.	2 2	4	-	8 4	Not available			<i>E. coli</i> <i>P. aeruginosa</i>	ATCC® 25922 ATCC® 27853	0.25-2 0.5-2
QUINUPRISTIN-DALFOPRISTIN 0.002 - 32 µg/mL	QDA	<i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>S. pneumoniae</i> <i>Streptococcus</i> spp. β-Hemolytic Group <i>Streptococcus</i> spp. Viridans Group	1 1 1 1 1	2	2	4 4 4 4 4	<i>Staphylococcus</i> spp. <i>Enterococcus</i> spp.	1 1	2 4	<i>S. aureus</i> <i>E. faecalis</i> <i>H. influenzae</i> <i>S. pneumoniae</i>	ATCC® 29213 ATCC® 29212 ATCC® 49247 ATCC® 49619	0.25-1 2-8 2-8 0.25-1

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
RIFAMPICIN 0.002 - 32 µg/mL 0.016 - 256 µg/mL	RD	<i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>Haemophilus</i> spp. <i>S. pneumoniae</i> <i>N. meningitidis</i>	1 1 1 1 0.5		2 2 2 2 1	4 4 4 4 2	<i>Staphylococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G <i>S. pneumoniae</i> <i>H. influenzae</i> <i>N. meningitidis</i> <i>H. pylori</i> <i>Corynebacterium</i> spp. <i>Aerococcus sanguinicola</i> and <i>urinae</i> <i>Kingella kingae</i>	0.06 0.06 0.06 1 0.25 1 0.06 0.12 0.5	0.5 0.5 0.5 1 0.25 1 0.5 0.12 0.5	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>H. influenzae</i> <i>S. pneumoniae</i> <i>H. influenzae</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 49247 ATCC® 49619 ATCC® 49766	0.004-0.016 0.5-4 4-16 16-64 0.25-1 0.016-0.06 0.25-1
SOLITHROMYCIN 0.002 - 32 µg/mL	SOL	Not available					Not available			<i>S. aureus</i> <i>E. faecalis</i> <i>H. influenzae</i> <i>S. pneumoniae</i>	ATCC® 29213 ATCC® 29212 ATCC® 49247 ATCC® 49619	0.03-0.12 0.016-0.06 1-4 0.004-0.016
SPECTINOMYCIN 0.064 - 1024 µg/mL	SPC	<i>N. gonorrhoeae</i>	32		64	128	<i>N. gonorrhoeae</i>	64	64	<i>N. gonorrhoeae</i>	ATCC® 49226	8-32
STREPTOMYCIN 0.016 - 256 µg/mL 0.064 - 1024 µg/mL	S	<i>Enterococcus</i> spp. (screening test for HLAR)	500		-	1000	<i>Enterococcus</i> spp. (test for high-level streptomycin resistance)	512	512	HLAR <i>E. faecalis</i> <i>E. faecalis</i>	ATCC® 29212 ATCC® 51299	< 500 ≥ 1000
TEICOPLANIN 0.016 - 256 µg/mL	TEC	<i>Staphylococcus</i> spp. <i>Enterococcus</i> spp.	8 8		16 16	32 32	<i>S. aureus</i> Coagulase-negative staphylococci <i>Enterococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G <i>S. pneumoniae</i> Viridans group streptococci	2 4 2 2 2 2	2 4 2 2 2 2	<i>S. aureus</i> <i>E. faecalis</i>	ATCC® 29213 ATCC® 29212	0.25-1 0.25-1
TEDIZOLID 0.002 - 32 µg/mL	TZD	<i>S. aureus</i> (including MRSA) <i>E. faecalis</i> <i>S. pyogenes</i> and <i>S. agalactiae</i> <i>S. anginosus</i> Group	0.5 0.25 0.5 0.25		1 - - -	2 - - -	<i>Staphylococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G <i>S. anginosus</i> Group	0.5 0.5 0.25	0.5 0.5 0.25	<i>S. aureus</i> <i>E. faecalis</i> <i>S. pneumoniae</i>	ATCC® 29213 ATCC® 29212 ATCC® 49619	0.12-1 0.25-1 0.12-0.5
TELAVANCIN 0.002 - 32 µg/mL 0.016 - 256 µg/mL	TLV	<i>S. aureus</i> (including MRSA) Vancomycin-susceptible <i>E. faecalis</i> <i>Streptococcus</i> spp. β-Hemolytic Group <i>Streptococcus</i> spp. Viridans Group	0.12 0.25 0.12 0.06		- - - -	- - - -	Methicillin-resistant <i>S. aureus</i>	0.125	0.125	<i>S. aureus</i> <i>E. faecalis</i> <i>S. pneumoniae</i>	ATCC® 29213 ATCC® 29212 ATCC® 49619	0.03-0.12 0.03-0.12 0.004-0.016
TETRACYCLINE 0.016 - 256 µg/mL	TE	Enterobacteriaceae <i>Acinetobacter</i> spp. Other Non-Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>Haemophilus</i> spp. <i>N. gonorrhoeae</i> <i>S. pneumoniae</i> <i>Streptococcus</i> spp. β-Hemolytic Group <i>Streptococcus</i> spp. Viridans Group Anaerobes	4 4 4 4 4 2 0.25 1 2 2 2 4		8 8 8 8 8 4 0.5-1 2 4 4 4 8	16 16 16 16 16 8 2 4 8 8 8 16	<i>Staphylococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G <i>S. pneumoniae</i> <i>H. influenzae</i> <i>M. catarrhalis</i> <i>N. gonorrhoeae</i> <i>N. meningitidis</i> <i>H. pylori</i> <i>C. jejuni</i> and <i>coli</i> <i>Corynebacterium</i> spp. <i>Kingella kingae</i>	1 1 1 1 1 0.5 1 1 2 2 2 0.5	2 2 2 2 2 1 1 2 2 2 2 0.5	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>H. influenzae</i> <i>S. pneumoniae</i> <i>N. gonorrhoeae</i> <i>B. fragilis</i> <i>B. thetaiotaomicron</i> <i>H. influenzae</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 49247 ATCC® 49619 ATCC® 49226 ATCC® 25285 ATCC® 29741 ATCC® 49766	0.12-1 8-32 0.5-2 8-32 4-32 0.06-0.5 0.25-1 0.125-0.5 8-32 0.25-1
TIAMULIN 0.002 - 32 µg/mL	TIA	Refer to CLSI VET01S					Not available			<i>S. aureus</i> <i>S. pneumoniae</i>	ATCC® 29213 ATCC® 49619	0.5-2 0.5-4

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL	MIC µg/mL	
TICARCILLIN 0.016 - 256 µg/mL	TC	Not available					Enterobacteriaceae <i>Pseudomonas</i> spp. Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes PK/PD (Non-species related) breakpoints	8 16 8 16 8	16 16 16 16 16	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>E. coli</i> <i>K. pneumoniae</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 35218 ATCC® 700603	2-8 16-64 4-16 8-32 > 128 > 256
TICARCILLIN*- CLAVULANIC ACID (2 µg/mL) 0.016 - 256* µg/mL	TTC	Enterobacteriaceae <i>P. aeruginosa</i> <i>Acinetobacter</i> spp. <i>B. cepacia</i> <i>S. maltophilia</i> Other Non-Enterobacteriaceae Anaerobes	16 16 16 16 16 16		32-64 32-64 32-64 32-64 32-64 32-64	128 128 128 128 128 128	Enterobacteriaceae <i>Pseudomonas</i> spp. Gram-positive anaerobes except <i>C. difficile</i> Gram-negative anaerobes PK/PD (Non-species related) breakpoints	8 16 8 8 8	16 16 16 16 16 16	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>E. coli</i> <i>K. pneumoniae</i> <i>B. fragilis</i> <i>B. thetaiotaomicron</i> <i>E. lenta</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 35218 ATCC® 700603 ATCC® 25285 ATCC® 29741 ATCC® 43055	0.5-2 16-64 4-16 8-32 8-32 32-128 0.06-0.5 0.5-2 8-32
TIGECYCLINE 0.016 - 256 µg/mL	TGC	Refer to the FDA drug label					Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G PK/PD (Non-species related) breakpoints	1 0.5 0.25 0.25 0.25	2 0.5 0.5 0.5 0.5	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>H. influenzae</i> <i>S. pneumoniae</i> <i>B. fragilis</i> <i>B. thetaiotaomicron</i> <i>C. difficile</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 49247 ATCC® 49619 ATCC® 25285 ATCC® 29741 ATCC® 700057	0.03-0.25 0.03-0.12 0.03-0.25 0.06-0.5 0.016-0.12 0.06-0.5 0.25-1 0.03-0.12
TILMICOSIN 0.002 - 32 µg/mL	TIL	Refer to CLSI VET01S					Not available			<i>S. aureus</i> <i>E. faecalis</i>	ATCC® 29213 ATCC® 29212	1-4 8-32
TOBRAMYCIN 0.016 - 256 µg/mL 0.064 - 1024 µg/mL	TOB	Enterobacteriaceae <i>P. aeruginosa</i> <i>Acinetobacter</i> spp. Other Non-Enterobacteriaceae <i>Staphylococcus</i> spp.	4 4 4 4 4		8 8 8 8 8	16 16 16 16 16	Enterobacteriaceae <i>Pseudomonas</i> spp. <i>Acinetobacter</i> spp. <i>S. aureus</i> Coagulase-negative staphylococci PK/PD (Non-species related) breakpoints	2 4 4 1 1 2	4 4 4 1 1 4	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853	0.12-1 8-32 0.25-1 0.25-1
TRIMETHOPRIM 0.002 - 32 µg/mL	TM	Enterobacteriaceae <i>Staphylococcus</i> spp.	8 8		- -	16 16	Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G	2 2 0.03 2	4 4 1 2	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853	1-4 0.12-0.5 0.5-2 > 64
TRIMETHOPRIM*- SULFAMETHOXAZOLE (1/19) 0.002 - 32* µg/mL	SXT	Enterobacteriaceae <i>Acinetobacter</i> spp. <i>B. cepacia</i> <i>S. maltophilia</i> Other Non-Enterobacteriaceae <i>Staphylococcus</i> spp. <i>Haemophilus</i> spp. <i>S. pneumoniae</i> <i>N. meningitidis</i>	2 2 2 2 2 2 0.5 0.5 0.12		- - - - - - 1-2 1-2 0.25	4 4 4 4 4 4 4 4 0.5	Enterobacteriaceae <i>S. maltophilia</i> <i>Acinetobacter</i> spp. <i>Staphylococcus</i> spp. <i>Enterococcus</i> spp. <i>Streptococcus</i> groups A, B, C and G <i>S. pneumoniae</i> <i>H. influenzae</i> <i>M. catarrhalis</i> <i>L. monocytogenes</i> <i>P. multocida</i> <i>Kingella kingae</i>	2 4 2 2 0.03 1 1 0.5 0.5 0.06 0.25 0.25	4 4 4 4 1 2 2 1 1 0.06 0.25 0.25	<i>S. aureus</i> <i>E. faecalis</i> <i>E. coli</i> <i>P. aeruginosa</i> <i>H. influenzae</i> <i>S. pneumoniae</i> <i>H. influenzae</i>	ATCC® 29213 ATCC® 29212 ATCC® 25922 ATCC® 27853 ATCC® 49247 ATCC® 49619 ATCC® 49766	≤ 0.5 ≤ 0.5 ≤ 0.5 8-32 0.03-0.25 0.12-1 0.016-0.06

ANTIBIOTIC	CODE	CLSI INTERPRETATIVE CRITERIA <sup>1</sup>	S≤	SDD MIC µg/mL	I	R≥	EUCAST INTERPRETATIVE CRITERIA <sup>2</sup>	S≤	R>	QUALITY CONTROL		
										MIC µg/mL	MIC µg/mL	
VANCOMYCIN 0.016 - 256 µg/mL	VA	<i>S. aureus</i>	2		4-8	16	<i>S. aureus</i>	2	2	<i>S. aureus</i>	ATCC® 29213	0.5-2
		<i>Staphylococcus</i> spp.	4		8-16	32	Coagulase-negative staphylococci	4	4	<i>E. faecalis</i>	ATCC® 29212	1-4
		<i>Enterococcus</i> spp.	4		8-16	32	<i>Enterococcus</i> spp.	4	4	<i>S. pneumoniae</i>	ATCC® 49619	0.12-0.5
		<i>S. pneumoniae</i>	1		-	-	<i>Streptococcus</i> groups A, B, C and G	2	2			
		<i>Streptococcus</i> spp. β-Hemolytic Group	1		-	-	<i>S. pneumoniae</i>	2	2			
		Viridans group streptococci					Viridans group streptococci	2	2			
		<i>Streptococcus</i> spp. Viridans Group	1		-	-	Gram-positive anaerobes except <i>C. difficile</i>	2	2			
							<i>C. difficile</i>	2	2			
							<i>Corynebacterium</i> spp.	2	2			
							<i>Aerococcus sanguinicola</i> and <i>urinae</i>	1	1			

\* Value on the M.I.C. scale refers to the first component of the combination.

ANTIFUNGAL	CODE	CLSI INTERPRETATIVE CRITERIA <sup>5</sup>			EUCAST INTERPRETATIVE CRITERIA <sup>6</sup>			QUALITY CONTROL			
		S≤	I	R≥	S≤	R>	MIC µg/mL	MIC µg/mL	MIC µg/mL	48 hours	
AMPHOTERICIN B 0.002 - 32 µg/mL	AMB	Not available			<i>C. albicans</i>	1	1	<i>C. parapsilosis</i>	ATCC® 22019	0.25-1	
					<i>C. glabrata</i>	1	1	<i>C. krusei</i>	ATCC® 6258	0.5-2	
					<i>C. krusei</i>	1	1	<i>C. albicans</i>	ATCC® 90028	0.125-0.5	
					<i>C. parapsilosis</i>	1	1				
					<i>C. tropicalis</i>	1	1				
					<i>A. fumigatus</i>	1	2				
					<i>A. niger</i>	1	2				
ANIDULAFUNGIN 0.002 - 32 µg/mL	AND	<i>C. albicans</i>	0.25	0.5	1	<i>C. albicans</i>	0.03	0.03	<i>C. parapsilosis</i>	ATCC® 22019	0.5-4
		<i>C. glabrata</i>	0.12	0.25	0.5	<i>C. glabrata</i>	0.06	0.06	<i>C. krusei</i>	ATCC® 6258	0.016-0.125
		<i>C. tropicalis</i>	0.25	0.5	1	<i>C. krusei</i>	0.06	0.06	<i>C. albicans</i>	ATCC® 90028	0.002-0.008
		<i>C. krusei</i>	0.25	0.5	1	<i>C. parapsilosis</i>	0.002	4			
		<i>C. parapsilosis</i>	2	4	8	<i>C. tropicalis</i>	0.06	0.06			
		<i>C. guilliermondii</i>	2	4	8						
CASPOFUNGIN 0.002 - 32 µg/mL	CAS	Not available			Not available			<i>C. parapsilosis</i>	ATCC® 22019	0.25-2	
								<i>C. krusei</i>	ATCC® 6258	0.25-1	
								<i>C. albicans</i>	ATCC® 90028	0.064-0.25	
FLUCONAZOLE 0.016 - 256 µg/mL	FLU	<i>C. albicans</i>	2	4	8	<i>C. albicans</i>	2	4	<i>C. parapsilosis</i>	ATCC® 22019	1-8
		<i>C. glabrata</i>	-	32	64	<i>C. glabrata</i>	0.002	32	<i>C. krusei</i>	ATCC® 6258	128≥256
		<i>C. parapsilosis</i>	2	4	8	<i>C. parapsilosis</i>	2	4	<i>C. albicans</i>	ATCC® 90028	0.125-0.5
		<i>C. tropicalis</i>	2	4	8	<i>C. tropicalis</i>	2	4			
FLUCYTOSINE 0.002 - 32 µg/mL	FC	Not available			Not available			<i>C. parapsilosis</i>	ATCC® 22019	0.064-0.25	
								<i>C. krusei</i>	ATCC® 6258	≥32	
								<i>C. albicans</i>	ATCC® 90028	0.5-2	
ITRACONAZOLE 0.002 - 32 µg/mL	ITC	Not available			<i>C. albicans</i>	0.06	0.06	<i>C. parapsilosis</i>	ATCC® 22019	0.064-0.25	
					<i>C. parapsilosis</i>	0.12	0.12	<i>C. krusei</i>	ATCC® 6258	0.25-1	
					<i>C. tropicalis</i>	0.12	0.12	<i>C. albicans</i>	ATCC® 90028	0.064-0.25	
					<i>A. flavus</i>	1	2				
					<i>A. fumigatus</i>	1	2				
					<i>A. nidulans</i>	1	2				
					<i>A. terreus</i>	1	2				
ISAVUCONAZOLE 0.002 - 32 µg/mL	IVU	Not available			<i>A. fumigatus</i>	1	1	<i>C. parapsilosis</i>	ATCC® 22019	0.008-0.032	
					<i>A. nidulans</i>	0.25	0.25	<i>C. krusei</i>	ATCC® 6258	0.125-1	
					<i>A. terreus</i>	1	1	<i>C. albicans</i>	ATCC® 90028	0.008-0.032	
KETOCONAZOLE 0.002 - 32 µg/mL	KE	Not available			Not available			<i>C. parapsilosis</i>	ATCC® 22019	0.032-0.125	
								<i>C. krusei</i>	ATCC® 6258	0.25-1	
								<i>C. albicans</i>	ATCC® 90028	0.008-0.032	
MICAFUNGIN 0.002 - 32 µg/mL	MYC	<i>C. albicans</i>	0.25	0.5	1	<i>C. albicans</i>	0.016	0.016	<i>C. parapsilosis</i>	ATCC® 22019	0.25-2
		<i>C. glabrata</i>	0.06	0.12	0.25	<i>C. glabrata</i>	0.03	0.03	<i>C. krusei</i>	ATCC® 6258	0.032-0.25
		<i>C. tropicalis</i>	0.25	0.5	1	<i>C. parapsilosis</i>	0.002	2	<i>C. albicans</i>	ATCC® 90028	0.004-0.032
		<i>C. krusei</i>	0.25	0.5	1						
		<i>C. parapsilosis</i>	2	4	8						
		<i>C. guilliermondii</i>	2	4	8						
POSACONAZOLE 0.002 - 32 µg/mL	POS	Not available			<i>C. albicans</i>	0.06	0.06	<i>C. parapsilosis</i>	ATCC® 22019	0.032-0.25	
					<i>C. parapsilosis</i>	0.06	0.06	<i>C. krusei</i>	ATCC® 6258	0.125-0.5	
					<i>C. tropicalis</i>	0.06	0.06	<i>C. albicans</i>	ATCC® 90028	0.032-0.125	
					<i>A. fumigatus</i>	0.12	0.25				
					<i>A. terreus</i>	0.12	0.25				
VORICONAZOLE 0.002 - 32 µg/mL	VO	<i>C. albicans</i>	0.12	0.25-0.5	1	<i>C. albicans</i>	0.12	0.12	<i>C. parapsilosis</i>	ATCC® 22019	0.016-0.064
		<i>C. krusei</i>	0.5	1	2	<i>C. parapsilosis</i>	0.12	0.12	<i>C. krusei</i>	ATCC® 6258	0.25-1
		<i>C. parapsilosis</i>	0.12	0.25-0.5	1	<i>C. tropicalis</i>	0.12	0.12	<i>C. albicans</i>	ATCC® 90028	0.004-0.016
		<i>C. tropicalis</i>	0.12	0.25-0.5	1	<i>A. fumigatus</i>	1	2			

ANTIMYCOBACTERIAL	CODE	INTERPRETATIVE CRITERIA	QUALITY CONTROL	MIC µg/mL
ETHAMBUTOL 0.016 - 256 µg/mL	EB	Not available	<i>M. tuberculosis</i>	0.064-0.25
ETHIONAMIDE 0.016 - 256 µg/mL	ET	Not available	<i>M. tuberculosis</i>	0.016-0.064
ISONIAZIDE 0.016 - 256 µg/mL	IZ	Not available	<i>M. tuberculosis</i>	0.016-0.064



## RESISTANCE DETECTION

ESBL: Extended Spectrum Beta Lactamase	CODE	INTERPRETATIVE CRITERIA <a href="#">Refer to MTS26</a>	QUALITY CONTROL	MIC µg/mL
CEFEPIME / CEFEPIME + CLAVULANIC ACID (4 µg/mL) 0.25-16 / 0.064-4 µg/mL	FEP/FEL	FEP ≥ 0.25 µg/mL and FEP/FEL ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>P. aeruginosa</i> ATCC® 27853 <i>K. pneumoniae</i> ATCC® 700603	FEP 0.5-2 FEL 1-4 FEP 0.25-1 FEL 0.064-0.25
CEFOTAXIME / CEFOTAXIME + CLAVULANIC ACID (4 µg/mL) 0.25-16 / 0.016-1 µg/mL	CTX/CTL	CTX ≥ 0.5 µg/mL and CTX/CTL ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>E. coli</i> ATCC® 35218 <i>K. pneumoniae</i> ATCC® 700603	CTX ≤ 0.25 CTL 0.016-0.064 CTX 1-4 CTL 0.125-1
CEFTAZIDIME / CEFTAZIDIME + CLAVULANIC ACID (4 µg/mL) 0.5-32 / 0.064-4 µg/mL	CAZ/CAL	CAZ ≥ 1 µg/mL and CAZ/CAL ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>E. coli</i> ATCC® 35218 <i>K. pneumoniae</i> ATCC® 700603	CAZ ≤ 0.5 CAL ≤ 0.064 CAZ ≥ 8 CAL 0.125-0.5

MBL: Metallo Beta Lactamase	CODE	INTERPRETATIVE CRITERIA <a href="#">Refer to MTS27</a>	QUALITY CONTROL	MIC µg/mL
IMIPENEM / IMIPENEM + EDTA 4-256 / 1-64 µg/mL 0.125-8 / 0.032-2 µg/mL	IMI/IMD	IMI/IMD ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>P. aeruginosa</i> ATCC® 27853 <i>S. maltophilia</i> ATCC® 13636	IMI ≤ 4 IMD 1-4 IMI 64-256 IMD 1-4
MEROPENEM / MEROPENEM + EDTA 0.125-8 / 0.032-2 µg/mL	MRP/MRD	MRP/MRD ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>P. aeruginosa</i> ATCC® 27853 <i>K. pneumoniae</i> ATCC® BAA-2146	MBL negative MBL positive

GRD: Glycopeptide Resistance Detection	CODE	INTERPRETATIVE CRITERIA <a href="#">Refer to MTS28</a>	QUALITY CONTROL	MIC µg/mL
VANCOMYCIN / TEICOPLANIN 0.5-32 / 0.5-32 µg/mL	VA/TEC	VA or TEC ≥ 8 µg/mL • GISA: GRD+ and <a href="#">MIC Test Strip VA</a> M.I.C. ≥ 4 µg/mL • hGISA: GRD+ and <a href="#">MIC Test Strip VA</a> M.I.C. < 4 µg/mL	<i>S. aureus</i> ATCC® 29213 <i>S. aureus</i> ATCC® 700698 <i>S. aureus</i> ATCC® 700699	VA 0.5-2 TEC 1-4 VA 1-8 TEC ≥ 32 VA 4-16 TEC ≥ 32

AmpC	CODE	INTERPRETATIVE CRITERIA <a href="#">Refer to MTS29</a>	Tentative QUALITY CONTROL	MIC µg/mL
CEFOTETAN / CEFOTETAN+ CLOXACILLIN 0.5-32 / 0.5-32 µg/mL	CTT/CXT	CTT/CXT ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>K. pneumoniae</i> ATCC® 700603 <i>K. pneumoniae</i> ATCC® BAA-1144	CTT 0.5-2 CXT 0.5-2 CTT ≥ 32 CXT 0.5-1

KPC: Klebsiella Producing Carbapenemase	CODE	INTERPRETATIVE CRITERIA <a href="#">Refer to MTS35</a>	QUALITY CONTROL	Result
ERTAPENEM / ERTAPENEM + PHENYLBORONIC ACID 0.125-8 / 0.032-2 µg/mL	ETP/EBO	ETP/EBO ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>K. pneumoniae</i> ATCC® BAA-1706	KPC negative
MEROPENEM / MEROPENEM + PHENYLBORONIC ACID 0.125-8 / 0.032-2 µg/mL	MRP/MBO	MRP/MBO ≥ 8 µg/mL or deformation of ellipse or phantom zone	<i>K. pneumoniae</i> ATCC® BAA-1705	KPC positive

**NB. These tests are NOT intended for the determination of the MIC.**

## References

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4. CLSI. Performance Standards for Antimicrobial Disk and Dilution Susceptibility Tests for Bacteria Isolated From Animals; Approved Standard - Third Edition. CLSI document M31-A3. Wayne, PA: Clinical and Laboratory Standards Institute; 2008.
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