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# Clindamycin Phosphate Susceptibility and Minimum Inhibitory Concentration (MIC) Data

## Microorganism Genus, Species, and Strain (if shown)

*Aggregatibacter actinomycetemcomitans*

*Bacillus circulans*

*Bacillus* spp.

*Bacillus* spp.

*Bacillus thetaiotaomicron*

*Bacteroides bivius*

*Bacteroides bivius*

*Bacteroides caccae*

*Bacteroides caccae*

*Bacteroides caccae*

*Bacteroides capillosus*

*Bacteroides capillosus*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis*

*Bacteroides distasonis* (ATCC 8503)

*Bacteroides eggerthii*

*Bacteroides eggerthii* (ATCC 27754)

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis*

*Bacteroides fragilis* (1995)

*Bacteroides fragilis* (1996)

*Bacteroides fragilis* (ATCC 23745)

*Bacteroides fragilis* (ATCC 25285)

*Bacteroides fragilis* (BF-1363 + clindamycin-resistant)

*Bacteroides fragilis* (GAI 0558)

*Bacteroides fragilis* (GAI 5562)

*Bacteroides fragilis* (GAI 7955)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (group)

*Bacteroides fragilis* (Imipenem-resistant)

*Bacteroides fragilis* (NCTC 10581)

*Bacteroides fragilis* (NCTC 9343)

*Bacteroides fragilis* (UC12199)

*Bacteroides fragilis* gr.

Concentration Range (µg/ml)

≤0.03 – >32

≤0.03 – >32

0.5 – >2

0.5 – >128

? – ?

≤0.062 – ?

≤0.5 – >125

≤0.06 – >256

0.25 – >128

≤0.03 – 2

0.5 – ?

0.03 – >128

≤0.5 – >128

≤0.06 – >256

≤0.062 – 4

0.5 – 8

≤0.06 – >256

0.03 – >32

0.03 – >16

≤0.03 – >128

0.5 – >64

0.06 – ?

≤0.5 – >128

≤0.03 – ?

0.03 – >128

≤0.004 – >128

≤0.03 – >128

? – ?

≤0.12 – >16

0.125 – 4

? – ?

≤0.5 – >128

0.06 – >32

0.125 – 8

0.125 – >16

0.25 – 4

0.125 – >128

≤0.06 – >256

0.125 – >256

0.5 – >32

0.5 – >8

0.015 – 2

0.5 – ?

0.5 – ?

0.5 – >256

≤0.25 – 256

0.06 – ?

1 – ?

>128 – ?

0.5 – ?

0.5 – ?

1 – ?

≤0.25 – >=512

0.01 – >=256

<0.125 – >128

≤0.016 – >32

≤0.5 – >128

≤0.016 – >32

≤0.06 – >256

0.008 – >64

0.06 – >128

1 – >128

≤0.03 – ?

0.5 – ?

1 – ?

≤0.06 – >256

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**Microorganism Genus, Species, and Strain (if shown)**

	Concentration Range (µg/ml)
<i>Bilophila wadsworthia</i>	0.25 – 2
<i>Bilophila wadsworthia</i> (WAL 7959)	0.5 – ?
<i>Brachyspira hyodysenteriae</i> (2000-2004 + Spain)	0.5 – >4
<i>Brachyspira hyodysenteriae</i> (2006-2007 + Spain)	0.5 – >4
<i>Brachyspira hyodysenteriae</i> (ATCC 27164)	0.06 – 0.125
<i>Brachyspira hyodysenteriae</i> (ATCC 27164)	0.063 – ?
<i>Brachyspira hyodysenteriae</i> (ATCC 27164)	≤0.125 – ?
<i>Brachyspira hyodysenteriae</i> (ATCC 31212)	1 – >4
<i>Brachyspira hyodysenteriae</i> (ATCC 31212)	>4 – ?
<i>Brevibacterium</i> spp.	0.5 – >64
<i>Callyspongia fallax</i>	0.02 – 8
<i>Campylobacter coli</i>	0.25 – >32
<i>Campylobacter concisus</i>	0.125 – 4
<i>Campylobacter gracilis</i>	0.125 – 4
<i>Campylobacter gracilis</i> (JCM 8538)	0.125 – ?
<i>Campylobacter jejuni</i>	0.12 – 8
<i>Campylobacter jejuni</i>	0.125 – 2
<i>Campylobacter mucosalis</i>	0.125 – 4
<i>Campylobacter rectus</i>	0.125 – 4
<i>Campylobacter showae</i>	0.125 – 4
<i>Campylobacter sputorum</i>	0.125 – 4
<i>Capnocytophaga ochracea</i>	≤0.03 – 2
<i>Capnocytophaga ochracea</i> (GAI 5586)	≤0.03 – ?
<i>Capnocytophaga</i> spp.	0.016 – >256
<i>Capnocytophaga</i> spp.	<1 – ?
<i>Cellulomonas biazotea</i>	2 – ?
<i>Cellulomonas cellasea</i>	0.06 – ?
<i>Cellulomonas fermentans</i>	0.25 – ?
<i>Cellulomonas fimi</i>	2 – ?
<i>Cellulomonas flavigena</i>	0.5 – ?
<i>Cellulomonas gelida</i>	1 – ?
<i>Cellulomonas hominis</i> (CDC A-3 + CE39)	8 – ?
<i>Cellulomonas hominis</i> (CDC A-3 + CE40)	8 – ?
<i>Cellulomonas uda</i>	2 – ?
<i>Clostridium baratii</i>	0.06 – >32
<i>Clostridium bifermentans</i>	0.02 – 8
<i>Clostridium bifermentans</i>	? – ?
<i>Clostridium bifermentans</i>	0.03 – 8
<i>Clostridium bifermentans</i>	0.016 – >64
<i>Clostridium bifermentans</i>	0.06 – >32
<i>Clostridium bifermentans</i>	0.06 – 0.25
<i>Clostridium bifermentans</i> (NCTC 06800)	1 – ?
<i>Clostridium butyricum</i>	0.02 – 8
<i>Clostridium butyricum</i>	≤0.03 – 4
<i>Clostridium cadaveris</i>	0.02 – 8
<i>Clostridium cadaveris</i>	0.03 – 8
<i>Clostridium cadaveris</i>	0.016 – >64
<i>Clostridium cadaveris</i>	0.06 – >32
<i>Clostridium cadaveris</i>	0.03 – ?
<i>Clostridium cadaveris</i>	0.5 – ?
<i>Clostridium clostridioforme</i>	≤0.015 – 2
<i>Clostridium clostridioforme</i>	0.02 – 2
<i>Clostridium clostridioforme</i>	≤0.03 – 4
<i>Clostridium clostridioforme</i>	0.06 – ?
<i>Clostridium clostridioforme</i> (NCTC 11224)	0.25 – ?
<i>Clostridium cochlearium</i>	0.02 – 8
<i>Clostridium difficile</i>	0.25 – >32
<i>Clostridium difficile</i>	? – ?
<i>Clostridium difficile</i>	1 – >32
<i>Clostridium difficile</i>	? – ?
<i>Clostridium difficile</i>	1 – >=32
<i>Clostridium difficile</i>	2 – >32
<i>Clostridium difficile</i>	2 – >32
<i>Clostridium difficile</i>	4 – >16
<i>Clostridium difficile</i>	8 – >16
<i>Clostridium difficile</i>	2 – >32
<i>Clostridium difficile</i>	4 – >64
<i>Clostridium difficile</i>	4 – >128
<i>Clostridium difficile</i>	4 – 128
<i>Clostridium difficile</i>	0.25 – 128
<i>Clostridium difficile</i>	6 – ?
<i>Clostridium difficile</i>	8 – ?

**Microorganism Genus, Species, and Strain (if shown)**

	<b>Concentration Range (µg/ml)</b>
<i>Clostridium difficile</i> (GAI 10029)	>128 – ?
<i>Clostridium glycolicum</i>	0.02 – 8
<i>Clostridium hastiforme</i>	0.06 – >32
<i>Clostridium histolyticum</i>	0.03 – 8
<i>Clostridium innocuum</i>	0.25 – 32
<i>Clostridium innocuum</i>	0.25 – >32
<i>Clostridium innocuum</i>	0.5 – 2
<i>Clostridium innocuum</i>	0.06 – >32
<i>Clostridium innocuum</i>	0.5 – 1
<i>Clostridium novyi</i> (A)	0.02 – 8
<i>Clostridium oroticum</i>	0.02 – 8
<i>Clostridium paraputrificum</i>	0.02 – 8
<i>Clostridium paraputrificum</i>	0.03 – 8
<i>Clostridium paraputrificum</i>	8 – ?
<i>Clostridium paraputrificum</i> (ATCC 25780)	8 – ?
<i>Clostridium perfringens</i>	0.008 – 4
<i>Clostridium perfringens</i>	≤0.016 – 2
<i>Clostridium perfringens</i>	0.06 – >32
<i>Clostridium perfringens</i>	? – ?
<i>Clostridium perfringens</i>	0.06 – 1
<i>Clostridium perfringens</i>	0.06 – 2
<i>Clostridium perfringens</i>	0.06 – 4
<i>Clostridium perfringens</i>	0.125 – 2
<i>Clostridium perfringens</i>	0.06 – 4
<i>Clostridium perfringens</i>	≤0.03 – >128
<i>Clostridium perfringens</i>	0.06 – 4
<i>Clostridium perfringens</i>	0.06 – >32
<i>Clostridium perfringens</i>	0.125 – >32
<i>Clostridium perfringens</i>	0.06 – 2
<i>Clostridium perfringens</i>	≤0.062 – 0.125
<i>Clostridium perfringens</i>	0.125 – 2
<i>Clostridium perfringens</i>	0.25 – ?
<i>Clostridium perfringens</i>	0.25 – ?
<i>Clostridium perfringens</i>	4.003 – ?
<i>Clostridium perfringens</i> (ATCC 13124)	0.06 – ?
<i>Clostridium putrificum</i> (ATCC 25784)	32 – ?
<i>Clostridium ramosum</i>	≤0.03 – 4
<i>Clostridium ramosum</i>	? – ?
<i>Clostridium ramosum</i>	0.25 – 4
<i>Clostridium ramosum</i>	0.06 – >32
<i>Clostridium ramosum</i>	0.5 – ?
<i>Clostridium ramosum</i>	6 – ?
<i>Clostridium ramosum</i> (ATCC 25582)	8 – ?
<i>Clostridium septicum</i>	≤0.03 – 4
<i>Clostridium septicum</i> (ATCC 12464)	≤0.03 – ?
<i>Clostridium sordellii</i>	0.03 – 8
<i>Clostridium sordellii</i>	0.016 – >64
<i>Clostridium sordellii</i>	0.06 – >32
<i>Clostridium sordellii</i>	1 – ?
<i>Clostridium sordellii</i> (NCTC 06929)	1 – ?
<i>Clostridium sordellii</i> (NCTC 08780)	1 – ?
<i>Clostridium species</i> (NOS)	? – ?
<i>Clostridium sporogenes</i>	≤0.03 – 4
<i>Clostridium sporogenes</i>	? – ?
<i>Clostridium spp.</i>	≤0.03 – 32
<i>Clostridium spp.</i>	0.008 – >128
<i>Clostridium spp.</i>	0.06 – 8
<i>Clostridium spp.</i>	0.03 – 8
<i>Clostridium spp.</i>	0.06 – >32
<i>Clostridium spp.</i>	0.016 – >64
<i>Clostridium spp.</i>	<0.03 – 4
<i>Clostridium tertium</i>	0.03 – 8
<i>Clostridium tertium</i>	0.016 – >64
<i>Clostridium tertium</i>	0.06 – >32
<i>Clostridium tertium</i>	1 – >32
<i>Collinsella aerofaciens</i>	≤0.03 – 0.25
<i>Corynebacterium</i> (CDC group G)	0.02 – >32
<i>Corynebacterium accolens</i>	0.02 – >32
<i>Corynebacterium amycolatum</i>	0.125 – >32
<i>Corynebacterium amycolatum</i>	0.5 – >64
<i>Corynebacterium aquaticum</i>	≤0.12 – >16
<i>Corynebacterium aquaticum</i>	0.02 – >32

**Microorganism Genus, Species, and Strain (if shown)**

	Concentration Range (µg/ml)
<i>Corynebacterium falsenii</i>	0.02 – >32
<i>Corynebacterium jeikeium</i>	≤0.12 – >16
<i>Corynebacterium jeikeium</i>	1 – >32
<i>Corynebacterium jeikeium</i>	0.5 – >64
<i>Corynebacterium minutissimum</i>	≤0.12 – >16
<i>Corynebacterium minutissimum</i>	0.02 – >32
<i>Corynebacterium minutissimum</i>	≤0.03 – >64
<i>Corynebacterium pseudodiphtheriticum</i>	0.02 – >32
<i>Corynebacterium</i> spp.	>8 – ?
<i>Corynebacterium</i> spp.	0.03 – >32
<i>Corynebacterium</i> spp.	0.12 – >128
<i>Corynebacterium</i> spp. (group 2)	≤0.12 – >16
<i>Corynebacterium</i> spp. (group ANF-1)	≤0.12 – >16
<i>Corynebacterium</i> spp. (group ANF-3)	≤0.12 – >16
<i>Corynebacterium striatum</i>	≤0.12 – >16
<i>Corynebacterium striatum</i>	0.02 – >32
<i>Corynebacterium striatum</i>	>64 – ?
<i>Corynebacterium urealyticum</i>	≤0.12 – >16
<i>Corynebacterium urealyticum</i>	0.02 – >32
<i>Corynebacterium urealyticum</i>	1 – >64
<i>Corynebacterium urealyticum</i> (with no species identified)	≤0.12 – >16
<i>Dermabacter hominis</i>	0.02 – >32
<i>Desulfovibrio piger</i> (DSM 749)	0.06 – ?
<i>Dialister pneumosintes</i>	≤0.03 – 2
<i>Edwardsiella hoshinae</i>	4 – 64
<i>Edwardsiella ictaluri</i>	1 – 32
<i>Edwardsiella tarda</i>	4 – 64
<i>Eggerthella lenta</i>	0.125 – 1
<i>Eggerthella lenta</i> (ATCC 25559)	0.25 – ?
<i>Eggerthella lenta</i> (ATCC 43055)	0.125 – ?
<i>Eikenella corrodens</i>	8 – >32
<i>Eikenella corrodens</i>	>64 – ?
<i>Enterococcus</i>	≤0.125 – 32
<i>Enterococcus avium</i>	≤0.12 – >16
<i>Enterococcus casseliflavus</i>	≤0.12 – >16
<i>Enterococcus cecorum</i>	≤0.12 – >16
<i>Enterococcus durans</i>	≤0.12 – >16
<i>Enterococcus faecalis</i>	0.016 – >256
<i>Enterococcus faecalis</i>	8 – >8
<i>Enterococcus faecalis</i>	≤0.12 – >16
<i>Enterococcus faecalis</i>	4 – ?
<i>Enterococcus faecalis</i> (ATCC 29212)	8 – ?
<i>Enterococcus faecalis</i> (erythromycin-resistant + vancomycin-susceptible)	16 – >256
<i>Enterococcus faecalis</i> (erythromycin-susceptible + vancomycin-susceptible)	16 – 32
<i>Enterococcus faecalis</i> (vancomycin-resistant)	>16 – ?
<i>Enterococcus faecalis</i> (vancomycin-resistant)	>256 – ?
<i>Enterococcus faecalis</i> (vancomycin-susceptible)	16 – >16
<i>Enterococcus faecalis</i> (vancomycin-susceptible)	16 – >256
<i>Enterococcus faecium</i>	0.12 – >8
<i>Enterococcus faecium</i> (12311 + vancomycin-resistant)	>128 – ?
<i>Enterococcus faecium</i> (12366 + vancomycin-resistant)	>128 – ?
<i>Enterococcus faecium</i> (ACA-DC 3350)	≤1 – ?
<i>Enterococcus faecium</i> (ACA-DC 3359)	≤1 – ?
<i>Enterococcus faecium</i> (erythromycin-resistant + vancomycin-resistant)	0.06 – >256
<i>Enterococcus faecium</i> (erythromycin-resistant + vancomycin-susceptible)	0.12 – >256
<i>Enterococcus faecium</i> (erythromycin-susceptible + vancomycin-susceptible)	0.12 – 16
<i>Enterococcus faecium</i> (PCD 71)	4 – ?
<i>Enterococcus faecium</i> (PCK 38)	2 – ?
<i>Enterococcus faecium</i> (PCK 45)	≤1 – ?
<i>Enterococcus faecium</i> (vancomycin-resistant)	≤0.12 – >16
<i>Enterococcus faecium</i> (vancomycin-resistant)	≤0.25 – >16
<i>Enterococcus faecium</i> (vancomycin-resistant)	? – ?
<i>Enterococcus faecium</i> (vancomycin-susceptible)	≤0.12 – >16
<i>Enterococcus faecium</i> (vancomycin-susceptible)	≤0.25 – >16
<i>Enterococcus faecium</i> (vancomycin-susceptible)	0.12 – >256
<i>Enterococcus faecium</i> (vancomycin-susceptible)	? – ?
<i>Enterococcus gallinarum</i>	≤0.12 – >16
<i>Enterococcus raffinosus</i>	≤0.12 – >16
<i>Enterococcus</i> spp.	0.12 – >8
<i>Enterococcus</i> spp.	8 – >16
<i>Enterococcus</i> spp. (macrolide-resistant)	>128 – ?
<i>Enterococcus</i> spp. (macrolide-susceptible)	0.25 – 32

**Microorganism Genus, Species, and Strain (if shown)**

	<b>Concentration Range (µg/ml)</b>
<i>Escherichia coli</i>	0.016 – >256
<i>Escherichia coli</i>	64 – ?
<i>Escherichia coli</i>	64 – ?
<i>Escherichia coli</i> (HB103::pBgIII)	≥128 – ?
<i>Escherichia coli</i> (HB103::pBluescript)	32 – ?
<i>Eubacterium brachy</i>	0.02 – 1
<i>Eubacterium combesii</i>	0.02 – 1
<i>Eubacterium contortum</i>	0.02 – 1
<i>Eubacterium lentum</i>	≤0.03 – 0.25
<i>Eubacterium lentum</i>	0.03 – 1
<i>Eubacterium lentum</i>	0.5 – 1
<i>Eubacterium lentum</i>	1 – ?
<i>Eubacterium lentum</i> (ATCC 43055)	0.02 – 1
<i>Eubacterium limosum</i>	0.02 – 1
<i>Eubacterium saburreum</i>	0.02 – 1
<i>Eubacterium saburreum</i>	≤0.03 – 0.25
<i>Eubacterium</i> spp.	<0.004 – 0.25
<i>Eubacterium</i> spp.	≤0.062 – 1
<i>Eubacterium</i> spp.	0.03 – 8
<i>Eubacterium</i> spp.	≤0.03 – 0.25
<i>Eubacterium</i> spp.	0.06 – 0.25
<i>Eubacterium tenue</i>	0.02 – 1
<i>Eubacterium timidum</i>	0.02 – 1
<i>Eubacterium timidum</i>	≤0.03 – 0.25
<i>Eubacterium yurii</i>	0.02 – 1
<i>Eubacterium yurii</i>	≤0.03 – 0.25
<i>Fingoldia magna</i>	≤0.03 – >32
<i>Fingoldia magna</i>	0.02 – 1
<i>Fingoldia magna</i>	0.125 – >32
<i>Fingoldia magna</i>	0.125 – >256
<i>Fingoldia magna</i>	0.125 – >16
<i>Fingoldia magna</i>	0.25 – 4
<i>Fingoldia magna</i> (ATCC 29328)	2 – ?
<i>Fingoldia magna</i> (Levofloxacin-resistant)	≤0.03 – >128
<i>Fusobacteria</i>	0.03 – 32
<i>Fusobacteria</i>	0.016 – 64
<i>Fusobacterium</i> (penicillin-susceptible)	0.03 – 0.25
<i>Fusobacterium</i> (penicillin-susceptible)	0.06 – 0.12
<i>Fusobacterium gonidiaformans</i>	≤0.015 – 2
<i>Fusobacterium mortiferum</i>	0.06 – 0.25
<i>Fusobacterium mortiferum</i>	0.06 – 8
<i>Fusobacterium mortiferum</i>	0.06 – 0.125
<i>Fusobacterium mortiferum</i>	0.125 – 4
<i>Fusobacterium mortiferum</i>	0.06 – 8
<i>Fusobacterium mortiferum</i>	0.25 – >128
<i>Fusobacterium mortiferum</i>	0.125 – ?
<i>Fusobacterium mortiferum</i>	18 – ?
<i>Fusobacterium naviforme</i>	≤0.03 – 0.06
<i>Fusobacterium naviforme</i>	≤0.015 – 2
<i>Fusobacterium necrogenes</i>	0.06 – 8
<i>Fusobacterium necrophorum</i>	≤0.03 – 0.06
<i>Fusobacterium necrophorum</i>	0.016 – 0.05
<i>Fusobacterium necrophorum</i>	0.03 – 0.06
<i>Fusobacterium necrophorum</i>	≤0.015 – 2
<i>Fusobacterium necrophorum</i>	0.03 – 0.125
<i>Fusobacterium necrophorum</i>	≤0.03 – 32
<i>Fusobacterium necrophorum</i>	0.25 – >128
<i>Fusobacterium necrophorum</i>	0.02 – 3.1
<i>Fusobacterium necrophorum</i>	0.03 – ?
<i>Fusobacterium necrophorum</i>	≤0.063 – ?
<i>Fusobacterium necrophorum</i> (ATCC 25286)	≤0.03 – ?
<i>Fusobacterium nucleatum</i>	? – ?
<i>Fusobacterium nucleatum</i>	≤0.03 – ≤0.03
<i>Fusobacterium nucleatum</i>	0.016 – >256
<i>Fusobacterium nucleatum</i>	0.06 – 0.125
<i>Fusobacterium nucleatum</i>	≤0.015 – 2
<i>Fusobacterium nucleatum</i>	0.03 – 0.125
<i>Fusobacterium nucleatum</i>	0.03 – 0.125
<i>Fusobacterium nucleatum</i>	0.06 – 0.125
<i>Fusobacterium nucleatum</i>	≤0.03 – 32
<i>Fusobacterium nucleatum</i>	0.25 – >128
<i>Fusobacterium nucleatum</i>	0.06 – 0.125

**Microorganism Genus, Species, and Strain (if shown)**

	<b>Concentration Range (µg/ml)</b>
<i>Fusobacterium nucleatum</i>	<=0.062 – 0.25
<i>Fusobacterium nucleatum</i>	<1 – ?
<i>Fusobacterium nucleatum</i> (ATCC 25586)	0.06 – ?
<i>Fusobacterium nucleatum</i> subsp. <i>animalis</i>	<=0.015 – 2
<i>Fusobacterium russii</i>	<=0.015 – 0.125
<i>Fusobacterium</i> spp.	0.015 – >128
<i>Fusobacterium</i> spp.	0.016 – >256
<i>Fusobacterium</i> spp.	0.06 – 8
<i>Fusobacterium</i> spp.	<=0.03 – 32
<i>Fusobacterium</i> spp.	0.03 – >32
<i>Fusobacterium</i> spp.	0.25 – >128
<i>Fusobacterium</i> spp.	0.25 – >128
<i>Fusobacterium ulcerans</i>	0.06 – 8
<i>Fusobacterium varium</i>	0.06 – 16
<i>Fusobacterium varium</i>	2 – >16
<i>Fusobacterium varium</i>	0.25 – >128
<i>Fusobacterium varium</i>	1 – >32
<i>Fusobacterium varium</i>	2 – 32
<i>Fusobacterium varium</i>	2 – 64
<i>Fusobacterium varium</i>	<=0.062 – 32
<i>Fusobacterium varium</i>	18 – ?
<i>Fusobacterium varium</i> (ATCC 8501)	4 – ?
<i>Gardnerella vaginalis</i>	≤0.03 – 0.25
<i>Gardnerella vaginalis</i> (ATCC 14018)	≤0.03 – ?
<i>Gemella morbillorum</i>	0.016 – >256
<i>Gemella morbillorum</i> (ATCC 27824)	0.06 – ?
<i>Gemella</i> spp.	≤0.12 – >16
<i>Haemophilus influenzae</i>	0.016 – >256
<i>Haemophilus influenzae</i>	0.5 – 16
<i>Haemophilus influenzae</i>	0.5 – 16
<i>Haemophilus influenzae</i> (ESBL)	1 – >4
<i>Haemophilus influenzae</i> (ESBL)	2 – >4
<i>Haemophilus influenzae</i> (ESBL)	2 – >4
<i>Haemophilus influenzae</i> (non-ESBL)	1 – >4
<i>Haemophilus parahaemolyticus</i>	? – ?
<i>Haemophilus parainfluenzae</i>	0.016 – >256
<i>Haemophilus parasuis</i> (Spain)	0.25 – 4
<i>Haemophilus parasuis</i> (UK)	0.25 – 4
<i>Helicobacter pylori</i>	0.1 – 50
<i>Hemolytic streptococci</i> (group C)	0.064 – >256
<i>Hemolytic streptococci</i> (group G)	0.064 – >256
<i>Jonesia denitrificans</i>	8 – ?
<i>Klebsiella pneumonia</i>	125 – ?
<i>Klebsiella pneumonia</i>	125 – ?
<i>Klebsiella pneumoniae</i>	0.016 – >256
<i>Lactobacillus acidophilus</i>	0.02 – >32
<i>Lactobacillus acidophilus</i>	≤0.125 – 0.25
<i>Lactobacillus acidophilus</i>	0.5 – 16
<i>Lactobacillus acidophilus</i> (CRL 1251)	0.1 – ?
<i>Lactobacillus acidophilus</i> (CRL 1266)	10 – ?
<i>Lactobacillus acidophilus</i> (Etest)	≤0.12 – >=256
<i>Lactobacillus acidophilus</i> (JCM 1132)	1 – ?
<i>Lactobacillus amylovorus</i> (Etest)	≤0.12 – >=256
<i>Lactobacillus brevis</i>	0.02 – >32
<i>Lactobacillus brevis</i>	4 – 32
<i>Lactobacillus brevis</i> (JCM 1059)	<=0.03 – ?
<i>Lactobacillus buchneri</i>	≤0.125 – 1
<i>Lactobacillus casei</i>	0.02 – >32
<i>Lactobacillus casei</i>	0.5 – 32
<i>Lactobacillus casei</i> (JCM 1134)	2 – ?
<i>Lactobacillus catenaforme</i>	≤0.03 – >32
<i>Lactobacillus catenaforme</i>	0.02 – >32
<i>Lactobacillus crispatus</i> (Etest)	≤0.12 – 2
<i>Lactobacillus curvatus</i>	8 – 64
<i>Lactobacillus delbrueckii</i>	≤0.03 – >32
<i>Lactobacillus delbrueckii</i>	0.25 – 64
<i>Lactobacillus delbrueckii</i> (Etest)	≤0.12 – 2
<i>Lactobacillus fermentans</i>	0.02 – >32
<i>Lactobacillus fermentum</i>	1 – 64
<i>Lactobacillus fermentum</i> (Etest)	≤0.12 – ?
<i>Lactobacillus fermentum</i> (JCM 1173)	<=0.03 – ?
<i>Lactobacillus gallinarum</i> (Etest)	≤0.12 – 1

**Microorganism Genus, Species, and Strain (if shown)**

	<b>Concentration Range (µg/ml)</b>
<i>Lactobacillus gasseri</i> (CRL 1259)	1 – ?
<i>Lactobacillus gasseri</i> (Etest)	≤0.12 – 16
<i>Lactobacillus helveticus</i> (Etest)	≤0.12 – 4
<i>Lactobacillus jensenii</i>	0.02 – >32
<i>Lactobacillus johnsonii</i> (CRL 1294)	>100 – ?
<i>Lactobacillus johnsonii</i> (Etest)	≤0.12 – 16
<i>Lactobacillus minutus</i>	0.02 – >32
<i>Lactobacillus oris</i>	≤0.03 – >32
<i>Lactobacillus oris</i>	0.02 – >32
<i>Lactobacillus paracasei</i> (CRL 1289)	>100 – ?
<i>Lactobacillus paracasei</i> (Etest)	≤0.25 – 4
<i>Lactobacillus paracasei</i> (MDIL)	≤0.25 – 16
<i>Lactobacillus pentosus</i> (PCD 101)	4 – ?
<i>Lactobacillus plantarum</i>	≤3 – >32
<i>Lactobacillus plantarum</i>	0.02 – >32
<i>Lactobacillus plantarum</i>	≤0.125 – >=256
<i>Lactobacillus plantarum</i>	0.5 – 128
<i>Lactobacillus plantarum</i> (Etest)	≤0.12 – 8
<i>Lactobacillus plantarum</i> (JCM 1149)	0.25 – ?
<i>Lactobacillus plantarum</i> (MDIL)	≤0.12 – 8
<i>Lactobacillus reuteri</i>	≤0.125 – 0.25
<i>Lactobacillus reuteri</i> (Etest)	≤0.12 – 2
<i>Lactobacillus reuteri</i> (JCM 1112)	<=0.03 – ?
<i>Lactobacillus reuteri</i> (MDIL)	≤0.12 – 16
<i>Lactobacillus rhamnosus</i>	0.02 – >32
<i>Lactobacillus rhamnosus</i>	≤0.125 – 0.5
<i>Lactobacillus rhamnosus</i>	0.5 – 32
<i>Lactobacillus rhamnosus</i> (Agar dilution)	≤0.125 – 8
<i>Lactobacillus rhamnosus</i> (Broth microdilution)	≤0.125 – 8
<i>Lactobacillus rhamnosus</i> (Etest)	≤0.12 – 16
<i>Lactobacillus rhamnosus</i> (Etest)	≤0.125 – 16
<i>Lactobacillus rhamnosus</i> (MDIL)	≤0.12 – 8
<i>Lactobacillus sakei</i>	8 – 64
<i>Lactobacillus sakei</i> (MDIL)	≤0.12 – 16
<i>Lactobacillus salivarius</i>	≤0.125 – 0.5
<i>Lactobacillus salivarius</i> (CRL 1328)	0.1 – ?
<i>Lactobacillus salivarius</i> (JCM 1231)	0.06 – ?
<i>Lactobacillus</i> sp.	0.25 – 128
<i>Lactobacillus</i> sp.	8 – ?
<i>Lactobacillus</i> spp.	≤0.03 – >32
<i>Lactobacillus</i> spp.	≤0.12 – >16
<i>Lactobacillus</i> spp.	0.015 – 128
<i>Lactobacillus</i> spp.	0.03 – 1
<i>Lactobacillus</i> spp.	0.25 – 4
<i>Lactobacillus</i> spp.	0.03 – 0.12
<i>Lactobacillus</i> spp.	0.5 – 1
<i>Lactobacillus</i> spp.	8 – ?
<i>Lactococcus</i>	≤0.125 – 32
<i>Leptotrichia buccalis</i>	≤0.03 – 0.06
<i>Leptotrichia buccalis</i>	0.125 – ?
<i>Leuconostoc</i>	≤0.125 – 32
<i>Leuconostoc mesenteroides</i> (PCD 119)	4 – ?
<i>Leuconostoc pseudomesenteroides</i> (PCK 18)	≤1 – ?
<i>Leuconostoc</i> spp.	≤0.12 – >16
<i>Leuconostoc</i> spp.	0.015 – 0.06
<i>Leuconostoc</i> spp.	<500 – ?
<i>Listeria monocytogenes</i>	≤0.12 – >16
<i>Listeria monocytogenes</i>	0.25 – 2
<i>Micrococcus</i> spp.	≤0.12 – >16
<i>Micromonas micros</i>	≤0.03 – 0.25
<i>Micromonas micros</i>	0.125 – 1
<i>Micromonas micros</i>	0.125 – 0.5
<i>Micromonas micros</i>	0.125 – 16
<i>Micromonas micros</i> (VPI 5464-1)	0.25 – ?
<i>Moraxella catarrhalis</i>	>2 – ?
<i>Moraxella catarrhalis</i>	1 – >4
<i>Moraxella catarrhalis</i>	1 – >4
<i>Moraxella catarrhalis</i>	0.016 – >256
<i>Mycobacterium smegmatis</i>	8 – ?
<i>Mycobacterium smegmatis</i> (A-2572U)	16 – ?
<i>Mycobacterium smegmatis</i> (C2055A)	16 – ?
<i>Mycobacterium smegmatis</i> (U2504G)	16 – ?



**Microorganism Genus, Species, and Strain (if shown)**

	<b>Concentration Range (µg/ml)</b>
<i>Mycoplasma fermentans</i>	≤0.008 – 0.031
<i>Mycoplasma fermentans</i>	≤0.008 – 0.063
<i>Mycoplasma hominis</i>	≤0.008 – 0.125
<i>Mycoplasma hominis</i>	≤0.008 – 0.125
<i>Mycoplasma hominis</i>	<2 – ?
<i>Mycoplasma hyopneumoniae</i> (F18)	0.125 – 0.25
<i>Mycoplasma hyopneumoniae</i> (F19)	>64 – ?
<i>Mycoplasma hyopneumoniae</i> (F2)	0.0625 – ?
<i>Mycoplasma hyopneumoniae</i> (F5)	0.125 – 0.25
<i>Mycoplasma hyopneumoniae</i> (F6)	0.125 – ?
<i>Mycoplasma hyopneumoniae</i> (J-strain)	0.5 – ?
<i>Mycoplasma hyosynoviae</i>	≤0.12 – 0.25
<i>Mycoplasma pneumonia</i>	? – ?
<i>Neisseria cinerea</i>	16 – >256
<i>Neisseria elongata</i>	16 – >256
<i>Neisseria gonorrhoeae</i>	0.5 – 4
<i>Neisseria meningitidis</i>	4 – ?
<i>Neisseria meningitidis</i>	4 – ?
<i>Neisseria sicca</i>	16 – >256
<i>Neisseria</i> spp.	16 – >256
<i>Nocardia asteroides</i>	100 – >=400
<i>Oerskovia turbata</i>	4 – ?
<i>Oerskovia xanthineolytica</i>	4 – ?
<i>Olsenella uli</i>	≤0.03 – >32
<i>Parvimonas micra</i>	0.047 – 2
<i>Pediococcus</i>	≤0.125 – 32
<i>Pediococcus pentosaceus</i> (PCD 215)	≤1 – ?
<i>Pediococcus pentosaceus</i> (PCD 237)	16 – ?
<i>Pediococcus</i> spp.	≤0.12 – >16
<i>Pediococcus</i> spp.	0.03 – 0.25
<i>Peptococcus asaccharolyticus</i>	? – ?
<i>Peptococcus asaccharolyticus</i>	≤0.062 – 1
<i>Peptococcus magnus</i>	≤0.062 – 2
<i>Peptococcus prevotii</i>	0.02 – 1
<i>Peptococcus</i> spp.	0.015 – >128
<i>Peptoniphilus asaccharolyticus</i>	0.06 – >16
<i>Peptoniphilus asaccharolyticus</i> (Levofloxacin-resistant)	≤0.03 – >128
<i>Peptoniphilus asaccharolyticus</i> (WAL 3218)	>128 – ?
<i>Peptoniphilus gorbachii</i>	0.125 – 0.75
<i>Peptoniphilus harei</i>	0.094 – 1.5
<i>Peptoniphilus indolicus</i> (GAI 0915)	32 – ?
<i>Peptoniphilus ivorii</i>	0.094 – 2
<i>Peptoniphilus lacrimalis</i>	0.016 – 0.38
<i>Peptoniphilus octavius</i>	0.047 – ?
<i>Peptostreptococcus</i> (penicillin-resistant)	≤0.015 – 2
<i>Peptostreptococcus</i> (penicillin-susceptible)	≤0.015 – 1
<i>Peptostreptococcus anaerobius</i>	0.03 – 0.5
<i>Peptostreptococcus anaerobius</i>	≤0.016 – 0.5
<i>Peptostreptococcus anaerobius</i>	≤0.03 – >32
<i>Peptostreptococcus anaerobius</i>	0.02 – 1
<i>Peptostreptococcus anaerobius</i>	≤0.016 – >32
<i>Peptostreptococcus anaerobius</i>	? – ?
<i>Peptostreptococcus anaerobius</i>	≤0.016 – >32
<i>Peptostreptococcus anaerobius</i>	0.015 – >128
<i>Peptostreptococcus anaerobius</i>	0.06 – 8
<i>Peptostreptococcus anaerobius</i>	0.06 – >128
<i>Peptostreptococcus anaerobius</i>	0.032 – 1
<i>Peptostreptococcus anaerobius</i>	≤0.062 – 0.125
<i>Peptostreptococcus anaerobius</i> (ATCC 27337)	0.25 – ?
<i>Peptostreptococcus anaerobius</i> (Levofloxacin-resistant)	≤0.03 – >128
<i>Peptostreptococcus asaccharolyticus</i>	≤0.015 – >32
<i>Peptostreptococcus asaccharolyticus</i>	0.02 – 1
<i>Peptostreptococcus asaccharolyticus</i>	0.06 – >32
<i>Peptostreptococcus asaccharolyticus</i>	≤0.016 – >32
<i>Peptostreptococcus asaccharolyticus</i>	≤0.016 – >32
<i>Peptostreptococcus asaccharolyticus</i>	0.06 – 8
<i>Peptostreptococcus magnus</i>	≤0.016 – >32
<i>Peptostreptococcus magnus</i>	≤0.016 – >32
<i>Peptostreptococcus magnus</i>	0.06 – 2
<i>Peptostreptococcus magnus</i>	0.06 – 8
<i>Peptostreptococcus magnus</i>	0.06 – >128
<i>Peptostreptococcus micros</i>	0.06 – 0.125

**Microorganism Genus, Species, and Strain (if shown)**

	Concentration Range (µg/ml)
<i>Peptostreptococcus micros</i>	<=0.016 - >32
<i>Peptostreptococcus micros</i>	<=0.016 - >32
<i>Peptostreptococcus micros</i>	0.06 - >128
<i>Peptostreptococcus micros</i>	<=0.062 - 0.5
<i>Peptostreptococcus prevotii</i>	<=0.016 - >32
<i>Peptostreptococcus prevotii</i>	<=0.016 - >32
<i>Peptostreptococcus prevotii</i>	0.03 - 32
<i>Peptostreptococcus prevotii</i>	0.125 - 0.25
<i>Peptostreptococcus sp.</i>	0.12 - 4
<i>Peptostreptococcus spp.</i>	0.008 - 64
<i>Peptostreptococcus spp.</i>	? - ?
<i>Peptostreptococcus spp.</i>	0.06 - >8
<i>Peptostreptococcus spp.</i>	0.125 - 256
<i>Peptostreptococcus spp.</i>	0.06 - >128
<i>Peptostreptococcus spp.</i>	0.06 - >128
<i>Peptostreptococcus spp.</i>	4 - ?
<i>Peptostreptococcus tetradius</i>	<=0.016 - >32
<i>Peptostreptococcus tetradius</i>	<=0.016 - >32
<i>Peptostreptococcus tetradius</i>	0.06 - 8
<i>Porphyromonas</i>	0.008 - 0.06
<i>Porphyromonas (penicillin-susceptible)</i>	≤0.015 - 0.06
<i>Porphyromonas asaccharolytica</i>	<=0.015 - >32
<i>Porphyromonas asaccharolytica</i>	0.008 - 0.03
<i>Porphyromonas asaccharolytica</i>	<=0.016 - >32
<i>Porphyromonas asaccharolytica</i>	<=0.016 - >32
<i>Porphyromonas asaccharolytica</i>	<=0.016 - ?
<i>Porphyromonas asaccharolytica (ATCC 25260)</i>	<=0.03 - ?
<i>Porphyromonas cangingivalis</i>	<=0.015 - ?
<i>Porphyromonas canoris</i>	<=0.015 - ?
<i>Porphyromonas cansulci</i>	<=0.015 - ?
<i>Porphyromonas circumdentaria</i>	<=0.015 - ?
<i>Porphyromonas endodontalis</i>	≤0.03 - 2
<i>Porphyromonas endodontalis</i>	0.016 - >256
<i>Porphyromonas gingivalis</i>	<=0.015 - ?
<i>Porphyromonas gingivalis</i>	<=0.016 - >32
<i>Porphyromonas gingivalis</i>	<=0.016 - >32
<i>Porphyromonas gingivalis</i>	≤0.03 - 2
<i>Porphyromonas gingivalis</i>	<=0.016 - ?
<i>Porphyromonas gingivalis</i>	<1 - ?
<i>Porphyromonas gingivalis (ATCC 33277)</i>	<=0.03 - ?
<i>Porphyromonas levii</i>	<=0.015 - ?
<i>Porphyromonas levii</i>	<=0.016 - ?
<i>Porphyromonas macacae</i>	<=0.015 - 0.03
<i>Porphyromonas spp.</i>	<=0.016 - >32
<i>Prevotella bivia</i>	<=0.015 - >32
<i>Prevotella bivia</i>	<=0.016 - >32
<i>Prevotella bivia</i>	<=0.016 - 0.06
<i>Prevotella bivia</i>	0.016 - 0.06
<i>Prevotella bivia</i>	<=0.016 - >16
<i>Prevotella bivia (ATCC 29303)</i>	<=0.03 - ?
<i>Prevotella buccae</i>	<=0.015 - 0.125
<i>Prevotella buccae</i>	0.008 - 0.03
<i>Prevotella buccae</i>	<=0.016 - >32
<i>Prevotella buccae</i>	<=0.016 - 0.03
<i>Prevotella buccae</i>	<=0.016 - >32
<i>Prevotella buccae</i>	0.016 - >256
<i>Prevotella buccae</i>	≤0.03 - >32
<i>Prevotella buccae</i>	<=0.03 - >128
<i>Prevotella buccae (ATCC 33574)</i>	<=0.03 - ?
<i>Prevotella buccalis</i>	≤0.03 - >32
<i>Prevotella corporis</i>	<=0.016 - >32
<i>Prevotella corporis</i>	<=0.016 - 0.125
<i>Prevotella corporis</i>	<=0.016 - 1
<i>Prevotella corporis</i>	0.016 - 0.03
<i>Prevotella corporis</i>	≤0.03 - <=0.03
<i>Prevotella corporis</i>	≤0.03 - >32
<i>Prevotella corporis (GAI 91000)</i>	<=0.03 - ?
<i>Prevotella dentalis</i>	≤0.03 - >32
<i>Prevotella denticola</i>	<=0.015 - 32
<i>Prevotella denticola</i>	0.016 - >256
<i>Prevotella denticola</i>	<=0.016 - >16
<i>Prevotella denticola</i>	≤0.03 - <=0.03

**Microorganism Genus, Species, and Strain (if shown)**

	<b>Concentration Range (µg/ml)</b>
<i>Prevotella denticola</i>	≤0.03 – >32
<i>Prevotella disiens</i>	≤0.016 – >32
<i>Prevotella disiens</i>	≤0.016 – >16
<i>Prevotella disiens</i>	≤0.016 – >32
<i>Prevotella disiens</i>	0.016 – ?
<i>Prevotella disiens</i>	≤0.03 – >32
<i>Prevotella heparinolytica</i>	≤0.015 – ?
<i>Prevotella heparinolytica</i> (ATCC 35895)	≤0.03 – ?
<i>Prevotella intermedia</i>	≤0.015 – 0.03
<i>Prevotella intermedia</i>	0.008 – 0.03
<i>Prevotella intermedia</i>	≤0.016 – 32
<i>Prevotella intermedia</i>	≤0.016 – ?
<i>Prevotella intermedia</i>	≤0.016 – >16
<i>Prevotella intermedia</i>	0.016 – >256
<i>Prevotella intermedia</i>	≤0.03 – >32
<i>Prevotella intermedia</i>	≤0.03 – >128
<i>Prevotella intermedia</i>	0.016 – ?
<i>Prevotella intermedia</i>	<1 – 1
<i>Prevotella intermedia</i> (ATCC 25611)	≤0.03 – ?
<i>Prevotella loescheii</i>	≤0.015 – 32
<i>Prevotella loescheii</i>	≤0.016 – >32
<i>Prevotella loescheii</i>	0.016 – >256
<i>Prevotella loescheii</i>	≤0.03 – ≤0.03
<i>Prevotella loescheii</i>	≤0.03 – >32
<i>Prevotella melaninogenica</i>	≤0.015 – 32
<i>Prevotella melaninogenica</i>	≤0.016 – >32
<i>Prevotella melaninogenica</i>	≤0.016 – 0.03
<i>Prevotella melaninogenica</i>	0.016 – >256
<i>Prevotella melaninogenica</i>	≤0.016 – >16
<i>Prevotella melaninogenica</i>	≤0.03 – ≤0.03
<i>Prevotella melaninogenica</i>	≤0.03 – >128
<i>Prevotella melaninogenica</i>	≤0.03 – >32
<i>Prevotella melaninogenica</i>	0.016 – 0.03
<i>Prevotella melaninogenica</i> (GAI 5490)	≤0.03 – ?
<i>Prevotella melaninogenicus</i>	≤0.03 – >128
<i>Prevotella nigrescens</i>	≤0.016 – >16
<i>Prevotella nigrescens</i>	≤0.03 – >32
<i>Prevotella oralis</i>	0.008 – 0.03
<i>Prevotella oralis</i>	≤0.016 – >32
<i>Prevotella oralis</i>	≤0.03 – >32
<i>Prevotella oralis</i>	0.016 – >256
<i>Prevotella oralis</i>	≤0.03 – >128
<i>Prevotella oralis</i> (ATCC 33269)	≤0.03 – ?
<i>Prevotella oralis</i> (group)	≤0.016 – >32
<i>Prevotella oris</i>	≤0.015 – 0.125
<i>Prevotella oris</i>	0.008 – 0.03
<i>Prevotella oris</i>	≤0.016 – >32
<i>Prevotella oris</i>	0.016 – >256
<i>Prevotella oris</i>	≤0.03 – >32
<i>Prevotella oris</i> (ATCC 33573)	≤0.03 – ?
<i>Prevotella pallens</i>	≤3 – >32
<i>Prevotella spp.</i>	? – ?
<i>Prevotella spp.</i>	≤0.016 – >32
<i>Prevotella spp.</i>	0.016 – >256
<i>Prevotella spp.</i>	≤0.03 – >32
<i>Prevotella spp.</i>	≤0.03 – >128
<i>Prevotella spp.</i>	≤0.03 – >8
<i>Prevotella spp.</i>	≤0.03 – >128
<i>Prevotella spp.</i>	<0.03 – >128
<i>Prevotella tanneræ</i>	≤0.03 – ≤0.03
<i>Prevotella tanneræ</i>	≤0.03 – >32
<i>Prevotella zoogloiformans</i>	≤0.03 – >32
<i>Propionibacteria</i>	0.06 – ?
<i>Propionibacterium</i>	? – ?
<i>Propionibacterium acnes</i>	≤0.03 – 0.25
<i>Propionibacterium acnes</i>	0.03 – 0.06
<i>Propionibacterium acnes</i>	0.06 – ?
<i>Propionibacterium acnes</i>	≤0.03 – 2
<i>Propionibacterium acnes</i>	0.06 – 0.5
<i>Propionibacterium acnes</i>	0.125 – 4
<i>Propionibacterium acnes</i>	0.01 – ?
<i>Propionibacterium acnes</i>	78 – ?

**Microorganism Genus, Species, and Strain (if shown)**

	Concentration Range (µg/ml)
<i>Propionibacterium acnes</i> (ATCC 11828)	0.125 – ?
<i>Propionibacterium avidum</i>	0.02 – 0.25
<i>Propionibacterium avidum</i>	≤0.0015 – 0.06
<i>Propionibacterium avidum</i>	≤0.03 – >32
<i>Propionibacterium granulosum</i>	≤0.015 – 0.03
<i>Propionibacterium granulosum</i>	0.02 – 0.25
<i>Propionibacterium granulosum</i> (ATCC 25564)	<=0.03 – ?
<i>Propionibacterium propionicus</i>	0.02 – 0.25
<i>Propionibacterium sp.</i>	0.016 – >256
<i>Propionibacterium spp.</i>	0.06 – 0.5
<i>Propionibacterium spp.</i>	0.125 – 256
<i>Proteus vulgaris</i>	250 – ?
<i>Proteus vulgaris</i>	250 – ?
<i>Pseudomonas aeruginosa</i>	1000 – ?
<i>Pseudomonas aeruginosa</i>	1000 – ?
<i>Rhodococcus spp.</i>	≤0.12 – >16
<i>Rhodococcus spp.</i>	0.25 – >256
<i>Rothia dentocariosa</i>	0.016 – >256
<i>Rothia mucilaginoso</i>	0.016 – >256
<i>Rothia spp.</i>	0.02 – >32
<i>Ruminococcus gnavus</i>	0.38 – ?
<i>Salmonella schottmuelleri</i>	64 – ?
<i>Salmonella schottmuelleri</i>	64 – ?
<i>Selenomonas flueggei</i>	≤0.03 – 2
<i>Selenomonas infelix</i>	≤0.03 – 2
<i>Selenomonas spp.</i>	≤0.03 – 2
<i>Staphylococci</i>	0.03 – 8
<i>Staphylococci</i>	0.5 – 8
<i>Staphylococci</i> (coagulase-negative + methicillin-resistant)	≤0.5 – >4
<i>Staphylococci</i> (coagulase-negative + methicillin-susceptible)	≤0.12 – >64
<i>Staphylococci</i> (coagulase-negative + methicillin-susceptible)	≤0.5 – >4
<i>Staphylococci</i> (coagulase-negative + oxacillin-resistant)	≤0.06 – >8
<i>Staphylococci</i> (coagulase-negative + oxacillin-resistant)	? – ?
<i>Staphylococci</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >16
<i>Staphylococci</i> (coagulase-negative + oxacillin-susceptible)	? – ?
<i>Staphylococci</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococci</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – 0.25
<i>Staphylococci</i> (coagulase-negative)	≤0.12 – >4
<i>Staphylococci</i> (coagulase-negative)	≤0.06 – >8
<i>Staphylococci</i> (coagulase-negative)	≤0.25 – >2
<i>Staphylococci</i> (coagulase-negative)	0.12 – >8
<i>Staphylococcus</i> (coagulase-negative + oxacillin-susceptible)	≤0.12 – >16
<i>Staphylococcus</i> (coagulase-negative + Uruguay)	≤1 – ?
<i>Staphylococcus</i> (coagulase-negative)	≤0.5 – >4
<i>Staphylococcus aureus</i>	≤0.12 – >4
<i>Staphylococcus aureus</i>	≤0.06 – >8
<i>Staphylococcus aureus</i>	0.03 – >16
<i>Staphylococcus aureus</i>	≤0.5 – >4
<i>Staphylococcus aureus</i>	0.016 – >256
<i>Staphylococcus aureus</i>	0.12 – >8
<i>Staphylococcus aureus</i>	? – ?
<i>Staphylococcus aureus</i>	0.04 – 1.6
<i>Staphylococcus aureus</i>	0.04 – 1.6
<i>Staphylococcus aureus</i>	43 – ?
<i>Staphylococcus aureus</i>	>64 – ?
<i>Staphylococcus aureus</i> (1-63)	0.06 – ?
<i>Staphylococcus aureus</i> (31593)	0.5 – ?
<i>Staphylococcus aureus</i> (ATCC 25923)	0.06 – ?
<i>Staphylococcus aureus</i> (ATCC 25923)	0.12 – ?
<i>Staphylococcus aureus</i> (ATCC 29213)	0.06 – ?
<i>Staphylococcus aureus</i> (ATCC 29213)	0.12 – ?
<i>Staphylococcus aureus</i> (ATCC 29213)	0.125 – ?
<i>Staphylococcus aureus</i> (ATCC 43300)	8192 – ?
<i>Staphylococcus aureus</i> (ATCC 6571)	0.06 – ?
<i>Staphylococcus aureus</i> (clindamycin-resistant + erythromycin-resistant + methicillin-resistant)	>128 – ?
<i>Staphylococcus aureus</i> (community-acquired + methicillin-resistant)	0.06 – 0.12
<i>Staphylococcus aureus</i> (community-acquired + methicillin-resistant)	? – ?
<i>Staphylococcus aureus</i> (community-acquired + methicillin-resistant)	≤0.25 – ?
<i>Staphylococcus aureus</i> (community-associated + methicillin-resistant)	≤0.25 – >8
<i>Staphylococcus aureus</i> (community-associated + methicillin-resistant)	≤0.12 – ?
<i>Staphylococcus aureus</i> (constitutively erm(A) + constitutively erm(C))	>128 – ?
<i>Staphylococcus aureus</i> (erythromycin-resistant + methicillin-resistant)	>256 – ?

**Microorganism Genus, Species, and Strain (if shown)**

	Concentration Range (µg/ml)
<i>Staphylococcus aureus</i> (erythromycin-resistant + methicillin-susceptible)	0.12 – ?
<i>Staphylococcus aureus</i> (erythromycin-resistant)	≤0.12 – >100
<i>Staphylococcus aureus</i> (erythromycin-susceptible + methicillin-susceptible)	0.12 – >256
<i>Staphylococcus aureus</i> (erythromycin-susceptible + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – ?
<i>Staphylococcus aureus</i> (health care-associated + methicillin-resistant)	≤0.25 – >8
<i>Staphylococcus aureus</i> (heterogeneous + vancomycin-intermediate)	≤0.12 – >4
<i>Staphylococcus aureus</i> (inducible erm(A))	0.06 – 0.12
<i>Staphylococcus aureus</i> (macrolide-susceptible)	0.06 – 0.12
<i>Staphylococcus aureus</i> (methicillin-resistant)	0.03 – >16
<i>Staphylococcus aureus</i> (methicillin-resistant)	≤0.5 – >4
<i>Staphylococcus aureus</i> (methicillin-resistant)	≤0.06 – >128
<i>Staphylococcus aureus</i> (methicillin-resistant)	? – ?
<i>Staphylococcus aureus</i> (methicillin-resistant)	64 – ?
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.06 – 0.12
<i>Staphylococcus aureus</i> (methicillin-susceptible)	0.06 – >16
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.12 – 16
<i>Staphylococcus aureus</i> (methicillin-susceptible)	0.12 – >256
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.5 – >4
<i>Staphylococcus aureus</i> (methicillin-susceptible)	? – ?
<i>Staphylococcus aureus</i> (methicillin-susceptible)	5 – ?
<i>Staphylococcus aureus</i> (methicillin-resistant)	≤0.12 – >4
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.12 – >4
<i>Staphylococcus aureus</i> (mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤0.25 – >2
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤? – >8
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤0.12 – >16
<i>Staphylococcus aureus</i> (oxacillin-resistant)	0.25 – >16
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus aureus</i> (oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus aureus</i> (erythromycin-resistant)	≤0.12 – >100
<i>Staphylococcus aureus</i> (erythromycin-susceptible + methicillin-susceptible)	0.12 – >256
<i>Staphylococcus aureus</i> (erythromycin-susceptible + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – ?
<i>Staphylococcus aureus</i> (health care-associated + methicillin-resistant)	≤0.25 – >8
<i>Staphylococcus aureus</i> (heterogeneous + vancomycin-intermediate)	≤0.12 – >4
<i>Staphylococcus aureus</i> (inducible erm(A))	0.06 – 0.12
<i>Staphylococcus aureus</i> (macrolide-susceptible)	0.06 – 0.12
<i>Staphylococcus aureus</i> (methicillin-resistant)	0.03 – >16
<i>Staphylococcus aureus</i> (methicillin-resistant)	≤0.5 – >4
<i>Staphylococcus aureus</i> (methicillin-resistant)	≤0.06 – >128
<i>Staphylococcus aureus</i> (methicillin-resistant)	? – ?
<i>Staphylococcus aureus</i> (methicillin-resistant)	64 – ?
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.06 – 0.12
<i>Staphylococcus aureus</i> (methicillin-susceptible)	0.06 – >16
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.12 – 16
<i>Staphylococcus aureus</i> (methicillin-susceptible)	0.12 – >256
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.5 – >4
<i>Staphylococcus aureus</i> (methicillin-susceptible)	? – ?
<i>Staphylococcus aureus</i> (methicillin-susceptible)	5 – ?
<i>Staphylococcus aureus</i> (methicillin-resistant)	≤0.12 – >4
<i>Staphylococcus aureus</i> (methicillin-susceptible)	≤0.12 – >4
<i>Staphylococcus aureus</i> (mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤0.25 – >2
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤? – >8
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤0.12 – >16
<i>Staphylococcus aureus</i> (oxacillin-resistant)	0.25 – >16
<i>Staphylococcus aureus</i> (oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus aureus</i> (oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus capitis</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus caprae</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus cohnii</i>	≤0.06 – >8
<i>Staphylococcus cohnii</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus epidermidis</i>	≤0.06 – >8
<i>Staphylococcus epidermidis</i>	0.016 – >256
<i>Staphylococcus epidermidis</i>	≤0.06 – >8
<i>Staphylococcus epidermidis</i>	0.1 – 0.2
<i>Staphylococcus epidermidis</i>	0.1 – 0.2
<i>Staphylococcus epidermidis</i>	62 – ?
<i>Staphylococcus epidermidis</i>	76 – ?
<i>Staphylococcus epidermidis</i> (coagulase-negative + mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus epidermidis</i> (coagulase-negative + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – >100
<i>Staphylococcus epidermidis</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus epidermidis</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8

**Microorganism Genus, Species, and Strain (if shown)****Concentration Range (µg/ml)**

<i>Staphylococcus epidermidis</i> (constitutively MLSB)	>128 – ?
<i>Staphylococcus epidermidis</i> (macrolide-susceptible)	0.06 – 0.12
<i>Staphylococcus epidermidis</i> (oxacillin-resistant)	≤0.5 – >4
<i>Staphylococcus epidermidis</i> (oxacillin-resistant)	≤0.25 – >16
<i>Staphylococcus epidermidis</i> (oxacillin-susceptible)	≤0.25 – >16
<i>Staphylococcus epidermidis</i> (oxacillin-susceptible)	≤0.5 – >4
<i>Staphylococcus haemolyticus</i>	≤0.06 – >8
<i>Staphylococcus haemolyticus</i>	≤0.25 – >2
<i>Staphylococcus haemolyticus</i>	0.12 – >8
<i>Staphylococcus haemolyticus</i> (coagulase-negative + mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus haemolyticus</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus haemolyticus</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus haemolyticus</i> (oxacillin-resistant)	≤0.25 – >16
<i>Staphylococcus haemolyticus</i> (oxacillin-susceptible)	≤0.25 – >16
<i>Staphylococcus hominis</i>	≤0.06 – >8
<i>Staphylococcus hominis</i> (coagulase-negative + mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus hominis</i> (coagulase-negative + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – >100
<i>Staphylococcus hominis</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus hominis</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus intermedius</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus intermedius</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus lugdunensis</i>	≤0.06 – >8
<i>Staphylococcus lugdunensis</i> (coagulase-negative + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – >100
<i>Staphylococcus lugdunensis</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus lugdunensis</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus pneumonia</i>	? – ?
<i>Staphylococcus pneumonia</i> (penicillin-resistant)	? – ?
<i>Staphylococcus pyogenes</i> (erythromycin-resistant)	≤0.12 – >100
<i>Staphylococcus saccharolyticus</i> (ATCC 14953)	0.125 – ?
<i>Staphylococcus saprophyticus</i>	≤0.06 – >8
<i>Staphylococcus saprophyticus</i> (coagulase-negative + mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus saprophyticus</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus saprophyticus</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus sciuri</i>	≤0.06 – >8
<i>Staphylococcus sciuri</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus simulans</i>	≤0.06 – >8
<i>Staphylococcus simulans</i> (coagulase-negative + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – >100
<i>Staphylococcus simulans</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus simulans</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus spp.</i>	≤0.06 – >8
<i>Staphylococcus spp.</i> (coagulase negative)	0.03 – >16
<i>Staphylococcus spp.</i> (coagulase negative)	? – ?
<i>Staphylococcus spp.</i> (coagulase-negative + methicillin-resistant)	≤0.06 – >128
<i>Staphylococcus spp.</i> (coagulase-negative + methicillin-susceptible)	≤0.06 – >128
<i>Staphylococcus spp.</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus spp.</i> (coagulase-negative)	≤0.25 – >16
<i>Staphylococcus warneri</i>	≤0.06 – >8
<i>Staphylococcus warneri</i> (coagulase-negative + mupirocin-resistant)	≤0.12 – >100
<i>Staphylococcus warneri</i> (coagulase-negative + mupirocin-susceptible + oxacillin-susceptible)	≤0.12 – >100
<i>Staphylococcus warneri</i> (coagulase-negative + oxacillin-resistant)	≤0.12 – >100
<i>Staphylococcus warneri</i> (coagulase-negative + oxacillin-susceptible)	≤0.06 – >8
<i>Staphylococcus xyloso</i>	≤0.06 – >8
<i>Stomatococcus spp.</i>	≤0.12 – >16
<i>Streptococci</i> (serogroup A + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup A + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup A)	0.06 – >256
<i>Streptococci</i> (serogroup B + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup B + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup B)	0.12 – ?
<i>Streptococci</i> (serogroup C + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup C + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup F + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup F + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup G + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (serogroup G + β-hemolytic)	≤0.06 – >8
<i>Streptococci</i> (Viridans group + penicillin-intermediate)	? – ?
<i>Streptococci</i> (Viridans group + penicillin-resistant)	0.5 – 8
<i>Streptococci</i> (Viridans group + penicillin-susceptible)	? – ?
<i>Streptococci</i> (Viridans group + penicillin-susceptible)	≤0.015 – 1
<i>Streptococci</i> (Viridans group)	≤0.03 – >0.25
<i>Streptococci</i> (Viridans group)	≤0.06 – >8
<i>Streptococci</i> (Viridans group)	≤0.12 – 4

**Microorganism Genus, Species, and Strain (if shown)**

	Concentration Range (µg/ml)
<i>Streptococci</i> ( $\alpha$ -hemolytic)	0.06 – 256
<i>Streptococci</i> ( $\beta$ -hemolytic)	$\leq 0.06$ – $> 8$
<i>Streptococci</i> ( $\beta$ -hemolytic)	$\leq 0.06$ – $> 8$
<i>Streptococci</i> ( $\beta$ -hemolytic)	$\leq 0.06$ – $> 8$
<i>Streptococci</i> ( $\beta$ -hemolytic)	? – ?
<i>Streptococci</i> ( $\beta$ -hemolytic)	$\leq 0.008$ – $> 16$
<i>Streptococcus</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus</i> (group A)	$\leq 0.03$ – $> 0.25$
<i>Streptococcus</i> (group B)	$\leq 0.03$ – $\geq 0.25$
<i>Streptococcus</i> (group C + erythromycin-susceptible)	$\leq 0.031$ – 0.125
<i>Streptococcus</i> (group C + <i>mef</i> (A))	0.063 – 0.125
<i>Streptococcus</i> (group G + erythromycin-susceptible)	$\leq 0.031$ – 0.25
<i>Streptococcus</i> (group G + inducible <i>erm</i> (TR))	0.063 – 0.5
<i>Streptococcus acidominimus</i>	0.064 – $> 256$
<i>Streptococcus agalactiae</i>	0.03 – $> 16$
<i>Streptococcus agalactiae</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus agalactiae</i>	0.03 – $> 64$
<i>Streptococcus agalactiae</i>	0.02 – 0.1
<i>Streptococcus agalactiae</i>	0.02 – 0.1
<i>Streptococcus agalactiae</i> (erythromycin-resistant)	0.03 – $> 64$
<i>Streptococcus agalactiae</i> (erythromycin-susceptible)	0.03 – 0.06
<i>Streptococcus anginosus</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus anginosus</i>	0.064 – $> 256$
<i>Streptococcus bovis</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus bovis</i>	$\leq 0.125$ – ?
<i>Streptococcus constellatus</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus constellatus</i>	0.064 – $> 256$
<i>Streptococcus constellatus</i> (ATCC 27923)	0.25 – ?
<i>Streptococcus dysgalactiae</i>	$\leq 0.06$ – $> 8$
<i>Streptococcus equisimilis</i>	$\leq 0.06$ – $> 8$
<i>Streptococcus infantarius</i>	$\leq 0.125$ – ?
<i>Streptococcus intermedius</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus intermedius</i>	$\leq 0.06$ – $> 8$
<i>Streptococcus intermedius</i>	0.064 – $> 256$
<i>Streptococcus intermedius</i> (ATCC 27335)	0.25 – ?
<i>Streptococcus milleri</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus mitis</i>	0.064 – $> 256$
<i>Streptococcus morbillorum</i>	0.5 – ?
<i>Streptococcus mutans</i>	$\leq 0.12$ – $> 16$
<i>Streptococcus oralis</i>	0.064 – $> 256$
<i>Streptococcus ovis</i>	0.064 – $> 256$
<i>Streptococcus parasanguinis</i>	0.064 – $> 256$
<i>Streptococcus plurimalium</i>	0.064 – $> 256$
<i>Streptococcus pneumonia</i>	0.015 – 0.25
<i>Streptococcus pneumonia</i>	$\leq 0.06$ – $> 8$
<i>Streptococcus pneumonia</i>	0.016 – 64
<i>Streptococcus pneumonia</i>	$\leq 0.12$ – 8
<i>Streptococcus pneumonia</i>	0.064 – $> 256$
<i>Streptococcus pneumonia</i>	$\leq 0.25$ – $> 16$
<i>Streptococcus pneumonia</i>	$\leq 0.25$ – $\geq 128$
<i>Streptococcus pneumonia</i>	$\leq 0.025$ – $> 1$
<i>Streptococcus pneumonia</i>	0.03 – $> 64$
<i>Streptococcus pneumonia</i>	0.002 – 0.1
<i>Streptococcus pneumonia</i>	0.002 – 0.1
<i>Streptococcus pneumonia</i>	0.125 – ?
<i>Streptococcus pneumonia</i>	0.125 – ?
<i>Streptococcus pneumonia</i>	0.25 – 128
<i>Streptococcus pneumonia</i> (ATCC 49619)	0.12 – ?
<i>Streptococcus pneumonia</i> (constitutively <i>erm</i> (B))	0.25 – $> 64$
<i>Streptococcus pneumonia</i> (Duke-2)	0.06 – ?
<i>Streptococcus pneumonia</i> ( <i>erm</i> (B) + <i>mef</i> (A))	0.12 – $> 16$
<i>Streptococcus pneumonia</i> ( <i>erm</i> (B) + <i>mef</i> (A))	0.12 – ?
<i>Streptococcus pneumonia</i> ( <i>erm</i> (B))	16 – $> 128$
<i>Streptococcus pneumonia</i> ( <i>erm</i> (B))	0.12 – $> 16$
<i>Streptococcus pneumonia</i> (erythromycin-susceptible)	$\leq 0.031$ – 0.063
<i>Streptococcus pneumonia</i> (macrolide-resistant)	41282
<i>Streptococcus pneumonia</i> (macrolide-susceptible)	0.008 – 0.06
<i>Streptococcus pneumonia</i> ( <i>mef</i> (A))	0.015 – 0.12
<i>Streptococcus pneumonia</i> ( <i>mef</i> (A))	0.063 – 0.125
<i>Streptococcus pneumonia</i> ( <i>mef</i> (A))	0.12 – 0.5
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	$\leq 0.03$ – $> 0.25$
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	$< 0.008$ – 8

**Microorganism Genus, Species, and Strain (if shown)****Concentration Range (µg/ml)**

<i>Streptococcus pneumonia</i> (penicillin-intermediate)	? – ?
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.12 – >16
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.12 – >16
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	<0.25 – >4
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.25 – >4
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	0.094 – >256
<i>Streptococcus pneumonia</i> (penicillin-resistant)	<0.008 – 8
<i>Streptococcus pneumonia</i> (penicillin-resistant)	≤0.06 – >8
<i>Streptococcus pneumonia</i> (penicillin-resistant)	? – ?
<i>Streptococcus pneumonia</i>	0.002 – 0.1
<i>Streptococcus pneumonia</i>	0.125 – ?
<i>Streptococcus pneumonia</i>	0.125 – ?
<i>Streptococcus pneumonia</i>	0.25 – 128
<i>Streptococcus pneumonia</i> (ATCC 49619)	0.12 – ?
<i>Streptococcus pneumonia</i> (constitutively erm(B))	0.25 – >64
<i>Streptococcus pneumonia</i> (Duke-2)	0.06 – ?
<i>Streptococcus pneumonia</i> (erm(B) + mef(A))	0.12 – >16
<i>Streptococcus pneumonia</i> (erm(B) + mef(A))	0.12 – ?
<i>Streptococcus pneumonia</i> (erm(B))	16 – >128
<i>Streptococcus pneumonia</i> (erm(B))	0.12 – >16
<i>Streptococcus pneumonia</i> (erythromycin-susceptible)	≤0.031 – 0.063
<i>Streptococcus pneumonia</i> (macrolide-susceptible)	0.008 – 0.06
<i>Streptococcus pneumonia</i> (mef(A))	0.015 – 0.12
<i>Streptococcus pneumonia</i> (mef(A))	0.063 – 0.125
<i>Streptococcus pneumonia</i> (mef(A))	0.12 – 0.5
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.03 – >0.25
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	<0.008 – 8
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	? – ?
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.12 – >16
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.12 – >16
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	<0.25 – >4
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	≤0.25 – >4
<i>Streptococcus pneumonia</i> (penicillin-intermediate)	0.094 – >256
<i>Streptococcus pneumonia</i> (penicillin-resistant)	<0.008 – 8
<i>Streptococcus pneumonia</i> (penicillin-resistant)	≤0.06 – >8
<i>Streptococcus pneumonia</i> (penicillin-resistant)	? – ?
<i>Streptococcus pyogenes</i> (erythromycin A-susceptible)	0.015 – 0.125
<i>Streptococcus pyogenes</i> (erythromycin-resistant)	0.1 – >64
<i>Streptococcus pyogenes</i> (erythromycin-susceptible)	0.03 – 0.06
<i>Streptococcus pyogenes</i> (erythromycin-susceptible)	≤0.031 – 0.125
<i>Streptococcus pyogenes</i> (inducible erm(A))	0.03 – 0.5
<i>Streptococcus pyogenes</i> (inducible erm(B) + mef(A))	0.25 – ?
<i>Streptococcus pyogenes</i> (inducible + erm(B))	0.25 – ?
<i>Streptococcus pyogenes</i> (inducible erm(TR) + mef(A))	0.063 – ?
<i>Streptococcus pyogenes</i> (inducible erm(TR))	0.063 – >64
<i>Streptococcus pyogenes</i> (macrolide-susceptible)	0.03 – 0.06
<i>Streptococcus pyogenes</i> (mef(A))	<0.015 – 0.06
<i>Streptococcus pyogenes</i> (mef(A))	≤0.031 – 0.125
<i>Streptococcus pyogenes</i> (mef(A))	? – 0.12
<i>Streptococcus salivarius</i>	0.064 – >256
<i>Streptococcus sanguinis</i>	0.064 – >256
<i>Streptococcus sobrinus</i>	0.064 – >256
<i>Streptococcus</i> spp.	≤0.06 – >8
<i>Streptococcus</i> spp. (serogroup A)	≤0.25 – ?
<i>Streptococcus</i> spp. (serogroup B)	≤0.25 – ?
<i>Streptococcus</i> spp. (serogroup C)	≤0.12 – >16
<i>Streptococcus</i> spp. (serogroup C)	≤0.25 – >8
<i>Streptococcus</i> spp. (serogroup F)	≤0.12 – >16
<i>Streptococcus</i> spp. (serogroup G)	≤0.12 – >16
<i>Streptococcus</i> spp. (serogroup G)	≤0.25 – >16
<i>Streptococcus</i> spp. (Viridans group)	≤0.12 – >16
<i>Streptococcus</i> spp. (Viridans group)	0.005 – 0.2
<i>Streptococcus</i> spp. (Viridans group)	0.005 – 0.2
<i>Streptococcus thoralensis</i>	0.064 – >256
<i>Sutterella wadsworthensis</i> (ATCC 51579)	16 – ?
<i>Veillonella dispar</i> (ATCC 17748)	<=0.03 – ?
<i>Veillonella parvula</i>	<=0.03 – >128
<i>Veillonella parvula</i>	0.125 – ?
<i>Veillonella parvula</i> (ATCC 10790)	<=0.03 – ?
<i>Veillonella</i> spp.	0.015 – <128
<i>Veillonella</i> spp.	≤0.03 – >32



**Microorganism Genus, Species, and Strain (if shown)**

*Veillonella* spp.

*Veillonella* spp.

*Veillonella* spp.

*Weissella* spp.

**Concentration Range (µg/ml)**

0.03 – >32

<=0.03 – >128

<0.03 – >128

≤0.125 – 1

The data above is sourced from The Antimicrobial Index. For further assistance, please contact us at [info@toku-e.com](mailto:info@toku-e.com) or visit [www.toku-e.com](http://www.toku-e.com).