# In Vitro Activity of Gemifloxacin (SB 265805) against Anaerobes

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Gemifloxacin mesylate (SB 265805), a new fluoronaphthyridone, was tested against 359 recent clinical anaerobic isolates by the National Committee for Clinical Laboratory Standards reference agar dilution method with supplemented brucella blood agar and an inoculum of 10<sup>5</sup> CFU/spot. Comparative antimicrobials tested included trovafloxacin, levofloxacin, grepafloxacin, sparfloxacin, sitafloxacin (DU-6859a), penicillin G, amoxicillin clavulanate, imipenem, cefoxitin, clindamycin, and metronidazole. The MIC<sub>50</sub> and MIC<sub>90</sub> (MICs at which 50 and 90% of the isolates were inhibited) of gemifloxacin against various organisms (with the number of strains tested in parentheses) were as follows (in micrograms per milliliter): for Bacteroides fragilis (28), 0.5 and 2; for Bacteroides thetaiotaomicron (24), 1 and 16; for Bacteroides caccae (12), 1 and 16; for Bacteroides distasonis (12), 8 and >16; for Bacteroides ovatus (12), 4 and >16; for Bacteroides stercoris (12), 0.5 and 0.5; for Bacteroides uniformis (12), 1 and 4; for Bacteroides vulgatus (11), 4 and 4; for Clostridium clostridioforme (15), 0.5 and 0.5; for Clostridium difficile (15), 1 and >16; for Clostridium innocuum (13), 0.125 and 2; for Clostridium perfringens (13), 0.06 and 0.06; for Clostridium ramosum (14), 0.25 and 8; for Fusobacterium nucleatum (12), 0.125 and 0.25; for Fusobacterium necrophorum (11), 0.25 and 0.5; for Fusobacterium varium (13), 0.5 and 1; for Fusobacterium spp. (12), 1 and 2; for Peptostreptococcus anaerobius (13), 0.06 and 0.06; for Peptostreptococcus asaccharolyticus (13), 0.125 and 0.125; for Peptostreptococcus magnus (14), 0.03 and 0.03; for Peptostreptococcus micros (12), 0.06 and 0.06; for Peptostreptococcus prevotii (14), 0.06 and 0.25; for Porphyromonas asaccharolytica (11), 0.125 and 0.125; for Prevotella bivia (10), 8 and 16; for Prevotella buccae (10), 2 and 2; for Prevotella intermedia (10), 0.5 and 0.5; and for Prevotella melaninogenica (11), 1 and 1. Gemifloxacin mesylate (SB 265805) was 1 to 4 dilutions more active than trovafloxacin against fusobacteria and peptostreptococci, and the two drugs were equivalent against clostridia and P. asaccharolytica. Gemifloxacin was equivalent to sitafloxacin (DU 6859a) against peptostreptococci, C. perfringens, and C. ramosum, and sitafloxacin was 2 to 3 dilutions more active against fusobacteria. Sparfloxacin, grepafloxacin, and levofloxacin were generally less active than gemifloxacin against all anaerobes.

Gemifloxacin mesylate (also called SB 265805 or LB20304), (R,S)-7-(3-aminomethyl-4-syn-methoxyimino-1-pyrrolidinyl)-1-cyclopropyl-6-fluoro-1,4-dihydro-4-oxo-1,8-naphthyridine-3carboxylic acid methanesulfonate, is a new fluoronaphthyridone with a broad spectrum of antimicrobial activity and enhanced activity against gram-positive aerobes (1, 3, 6). It is a racemic mixture (specific rotation = 0.0) with equipotent enantiomers (6a). Reports of several studies using a limited number of genera and strains have noted that it has potential activity against anaerobic bacteria, both gram positive and gram negative (1, 4). In order to further evaluate gemifloxacin's potential therapeutic utility against infections caused by anaerobic bacteria, we studied its in vitro activity against 359 recent clinical anaerobic isolates.

#### MATERIALS AND METHODS

The 359 anaerobic strains used had been isolated recently (from 1995 through 1998) from humans with clinical infections and identified by standard criteria (2, 7). The control strains *Bacteroides fragilis* ATCC 25285 and *Bacteroides thetaiotaonicron* ATCC 29741 were also included on each set of plates tested. Quality control was assured when limits approved by the National Committee for Clinical Laboratory Standards (NCCLS), were recorded for the various established compounds for *B. fragilis* ATCC 25285 and *B. thetaiotaomicron* ATCC 29741. *Fuso-*

*bacterium necrogenes* ATCC 25556 was included for comparative purposes. The numbers and species of isolates tested are given in Table 1.

Standard laboratory powders were supplied as follows: gemifloxacin and amoxicillin clavulanate by SmithKline Beecham, Philadelphia, Pa.; trovafloxacin by Pfizer Inc., New York, N.Y.; levofloxacin by R. W. Johnson Pharmaceutical Research Institute, Raritan, N.J.; grepafloxacin by Glaxo-Wellcome Inc., Research Triangle Park, N.C.; sparfloxacin by Rhone-Poulenc Rorer, Collegeville, Pa.; sitafloxacin by Daichi Pharmaceuticals, Tokyo, Japan; cefoxitin and imipenem by Merck & Co., West Point, Pa.; clindamycin by Pharmacia Upjohn Co., Kalamazoo, Mich.; metronidazole by Searle Research & Development, Skokie, III.; and penicillin G by Sigma Chemical Co., St. Louis, Mo.

Frozen cultures were transferred twice on brucella agar supplemented with hemin, vitamin  $K_1$ , and 5% sheep blood. Susceptibility testing was performed according to NCCLS standards (5). Brucella agar supplemented with hemin, vitamin  $K_1$ , and 5% laked sheep blood was the basal medium. Antimicrobial agents were reconstituted according to the manufacturers' instructions. Serial twofold dilutions of various concentrations of antimicrobial agents were prepared on the day of the test and added to the test agar medium.

The agar plates were inoculated with a Steers replicator (Craft Machine Inc., Chester, Pa.). The inoculum used was 10<sup>5</sup> CFU per spot. Plates were incubated in an anaerobic chamber for 48 h at 37°C prior to examination. The MIC was defined as the lowest concentration of an agent that yielded either no growth or a marked change in the appearance of growth compared to that on the growth control plate.

## **RESULTS AND DISCUSSION**

The comparative activities of gemifloxacin and the other agents tested are presented in Table 1. Our comparison of compounds was conducted on a MIC basis, since breakpoints have not yet been established for all of the compounds.

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TABLE 1. In vitro activity of gemifloxacin (SB 265805) compared to those of 11 other antimicrobial agents against 359	
TABLE 1. In vitro activity of geninioxacin (SB 205005) compared to those of 11 other antimicrobial agents against 559	
recently isolated clinical anaerobic bacterial strains	
recently isolated ennear anaeroole bacterial strains	

Organism (no. of isolates) and agent		MIC		Organism (no. of isolates)		MIC	
	Range	50%	90%	and agent	Range	50%	90%
Bacteroides fragilis (28)				Imipenem	0.125–1	0.25	0.5
Gemifloxacin	0.5-2	0.5	2	Metronidazole	1-4	2	2
Trovafloxacin	0.125-4	0.25	0.5	Clindamycin	1->128	2	>128
Sitafloxacin	0.06-0.25	0.06	0.25		1 - 120	-	- 120
Grepafloxacin	1–16	2	4	Bacteroides stercoris (12)			
Levofloxacin	1->16	1	4	Gemifloxacin	0.25-8	0.5	0.5
Sparfloxacin	0.5–16	1	2	Trovafloxacin	0.25-1	0.25	0.5
Penicillin G	0.5 -> 64	16	>64	Sitafloxacin	0.06-1	0.06	0.12
Amoxicillin clavulanate	0.25-32	0.5	8	Grepafloxacin	2-8	2	2
Cefoxitin	2-128	8	32	Levofloxacin	1-8	1	2
Imipenem	0.03 -> 32	0.125	2	Sparfloxacin	0.5-2	1	2
Metronidazole	0.05 > 52 0.5 > 16	2	$\frac{2}{2}$	Penicillin G	2->64	8	16
Clindamycin	$\leq 0.06 - > 128$	1	>128	Amoxicillin clavulanate	0.25-4	1	2
Cilildaniyeni	≤0.00->120	1	/120	Cefoxitin	1-32	8	32
acteroides thetaiotaomicron (24)					$\leq 0.015 - 0.5$	o 0.125	0.25
Gemifloxacin	0.5->16	1	16	Imipenem			
Trovafloxacin	0.25-4	0.5	4	Metronidazole	0.25-4	4	4
	0.23-4 0.06-16	0.3	2	Clindamycin	≤0.06->128	2	4
Sitafloxacin				Rastanoidas uniformis (12)			
Grepafloxacin	2->16	2	>16	Bacteroides uniformis (12)	0.125.16	1	4
Levofloxacin	2->16	4	>16	Gemifloxacin	0.125–16	1	4
Sparfloxacin	2->16	2	>16	Trovafloxacin	0.06-4	0.5	4
Penicillin G	4->64	32	>64	Sitafloxacin	0.03-4	0.125	4
Amoxicillin clavulanate	0.5 - 16	1	4	Grepafloxacin	0.25->16	4	8
Cefoxitin	4-128	32	64	Levofloxacin	≤0.5->16	4	>16
Imipenem	0.125-8	0.25	1	Sparfloxacin	0.25 -> 16	2	8
Metronidazole	0.5-4	2	2	Penicillin G	0.125 -> 64	16	>64
Clindamycin	$\leq 0.06 -> 128$	16	>128	Amoxicillin clavulanate	0.125-8	0.5	4
				Cefoxitin	1-128	8	32
Bacteroides caccae (12)				Imipenem	0.06 - 1	0.25	1
Gemifloxacin	0.5 - 16	1	16	Metronidazole	≤0.06–4	4	4
Trovafloxacin	0.125-4	0.5	2	Clindamycin	≤0.06->128	0.5	>128
Sitafloxacin	0.06 - 2	0.125	2				
Grepafloxacin	2->16	4	8	Bacteroides vulgatus (1)			
Levofloxacin	1->16	4	>16	Gemifloxacin	0.5-16	4	4
Sparfloxacin	0.5 - > 16	2	16	Trovafloxacin	0.125-4	0.25	4
Penicillin G	8-64	8	32	Sitafloxacin	0.06-4	0.125	1
Amoxicillin clavulanate	0.25-8	0.25	2	Grepafloxacin	2->16	4	8
Cefoxitin	8-32	16	16	Levofloxacin	1->16	2	>16
Imipenem	0.03-0.5	0.06	0.25	Sparfloxacin	0.5->16	1	16
Metronidazole	1-8	2	4	Penicillin G	1->64	64	>64
Clindamycin	0.25->128	$\frac{2}{2}$	>128	Amoxicillin clavulanate	0.25-8	2	2
Chindaniyeni	0.25->120	4	> 120	Cefoxitin	2-128	16	32
Bacteroides distasonis (12)				Imipenem	0.03-1	0.25	1
Gemifloxacin	0.5->16	8	>16	Metronidazole	1-4	2	2
Trovafloxacin	0.125-4	0.5	2			0.25	>128
Sitafloxacin	0.125-4	0.5	2	Clindamycin	≤0.06->128	0.23	~120
Grepafloxacin	1 - > 16	0.3 8	>16	<i>Clostridium clostridioforme</i> (15)			
Levofloxacin	1 > 10 1 - > 16	4	16	Gemifloxacin	0.125-1	0.5	0.5
Sparfloxacin	1-16	4	16	Trovafloxacin	0.125-1	4	8
						0.25	。 0.25
Penicillin G	8->64	16	>64	Sitafloxacin	0.125–0.5 1–16		
Amoxicillin clavulanate	0.25-8	2	8	Grepafloxacin		8	16
Cefoxitin	8-128	16	32	Levofloxacin	8 - > 16	8	>16
Imipenem	0.03-1	0.25	0.5	Sparfloxacin	4->16	8	16
Metronidazole	1-8	4	4	Penicillin G	0.5->32	2	32
Clindamycin	0.125->128	2	>128	Amoxicillin clavulanate	0.5-4	0.5	4
				Cefoxitin	2-16	4	16
acteroides ovatus (12)	0.7			Imipenem	0.06-2	1	2
Gemifloxacin	0.5->16	4	>16	Metronidazole	$\leq 0.06 - 0.5$	0.125	0.5
Trovafloxacin	0.5-8	1	4	Clindamycin	0.03-8	0.06	4
Sitafloxacin	0.125 -> 16	0.5	4				
Grepafloxacin	2->16	8	>16	Clostridium difficile (15)			
Levofloxacin	4->16	4	>16	Gemifloxacin	1->16	1	>16
Sparfloxacin	1->16	2	16	Trovafloxacin	0.5->16	1	>16
Penicillin G	16->64	32	>64	Sitafloxacin	0.125-1	0.25	1
Amoxicillin clavulanate	0.5-8	0.5	2	Grepafloxacin	4->16	16	16
Cefoxitin	16->128	32	128	Levofloxacin	$2 \rightarrow 10$ $2 \rightarrow 16$	4	>16

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TABLE 1-Continued

Organism (no. of isolates) and agent	MIC			Organism (no. of isolates)	MIC		
	Range	50%	90%	and agent	Range	50%	90%
Sparfloxacin	4->16	4	>16	Trovafloxacin	0.125-2	0.5	1
Penicillin G	1-4	2	4	Sitafloxacin	$\leq 0.015 - 0.06$	0.03	0.06
Amoxicillin clavulanate	0.5 - 1	1	1	Grepafloxacin	2-8	4	8
Cefoxitin		>32	>32	Levofloxacin	0.5-4	2	2
Imipenem	2–16	4	4	Sparfloxacin	0.5-4	1	2
Metronidazole	0.125–1	0.5	1	Penicillin G	≤0.015	≤0.015	≤0.015
Clindamycin	0.5 -> 128	3 4	>128	Amoxicillin clavulanate	$\leq 0.015 - 0.03$	$\leq 0.015$	0.03
				Cefoxitin	≤0.015-0.06	0.03	0.03
Clostridium innocuum (13)	0.06.2	0.495		Imipenem	≤0.015	≤0.015	≤0.015
Gemifloxacin	0.06-2	0.125	2	Metronidazole	$\leq 0.06 - 0.5$	0.125	0.5
Trovafloxacin	0.125–8 0.125–4	0.25	4	Clindamycin	≤0.015-0.03	≤0.015	0.03
Sitafloxacin Grepafloxacin	0.125-4 0.5->16	0.125	1 > 16	Fusobacterium varium (13)			
Levofloxacin	0.3 = >10 1 = >16	1 2	>10 16	Gemifloxacin	0.125-16	0.5	1
Sparfloxacin	0.25–16	0.5	16	Trovafloxacin	0.125 10	4	4
Penicillin G	0.25-0.5	0.25	0.5	Sitafloxacin	0.06-8	0.25	0.5
Amoxicillin clavulanate	0.125-0.5	0.25	0.5	Grepafloxacin	4->16	16	>16
Cefoxitin	8->32		>32	Levofloxacin	1->16	8	>16
Imipenem	0.5-2	1	2	Sparfloxacin	1->16	8	16
Metronidazole	0.25-1	0.5	0.5	Penicillin G	0.06 - 1	0.5	0.5
Clindamycin	0.25->128		>128	Amoxicillin clavulanate	0.25 - 2	1	2
5				Cefoxitin	0.25-8	4	8
Clostridium perfringens (13)				Imipenem	0.06-2	0.5	1
Gemifloxacin	0.03-0.125	0.06	0.06	Metronidazole	0.125-2	0.5	2
Trovafloxacin	0.06-0.25	0.125	0.25	Clindamycin	0.125-16	4	16
Sitafloxacin	≤0.015-0.125		0.06	<i>Fusobacterium</i> spp. (12) <sup>a</sup>			
Grepafloxacin	0.06-2	0.25	0.5	Gemifloxacin	0.125-8	1	2
Levofloxacin	0.25-1	0.25	1	Trovafloxacin	0.125 0	2	4
Sparfloxacin	0.03-0.5	0.125	0.25	Sitafloxacin	0.06-1	0.5	0.5
Penicillin G Amoxicillin clavulanate	0.03-0.125	$0.06 \le 0.015$	0.125 0.06	Grepafloxacin	8->16	16	>16
Cefoxitin	$\leq 0.015 - 0.06$ 0.25 - 2	$\leq 0.013$ 0.5	1	Levofloxacin	1->16	8	>16
Imipenem	0.03-0.125		0.06	Sparfloxacin	1->16	8	> 16
Metronidazole	0.5-2	1	2	Penicillin G	$\leq 0.015 -> 32$	0.25	1
Clindamycin	0.03-4	1	2	Amoxicillin clavulanate	0.5-32	1	4
				Cefoxitin	1-8	4	8
Clostridium ramosum (14)				Imipenem	0.25-1	0.5	1
Gemifloxacin	0.125-8	0.25	8	Metronidazole	0.25-2	0.5	1
Trovafloxacin	0.25 - 16	0.5	8	Clindamycin	0.06–32	0.125	32
Sitafloxacin	0.125-8	0.25	4	Peptostreptococcus anaerobius (13)			
Grepafloxacin	1->16	2	>16	Gemifloxacin	≤0.015-8	0.06	0.06
Levofloxacin	2->16	4	>16	Trovafloxacin	≤0.015-8	0.125	0.125
Sparfloxacin	0.5->16	2	>16	Sitafloxacin	≤0.015-0.5	0.03	0.03
Penicillin G	0.06-0.5	0.125	0.5	Grepafloxacin	0.25-8	1	1
Amoxicillin clavulanate	0.06-0.25	0.06	0.125	Levofloxacin	0.25 -> 16	0.25	0.5
Cefoxitin	4->32 0.125-0.5	8 0.25	>32 0.25	Sparfloxacin	0.125 -> 16	0.5	1
Imipenem Metronidazole	0.123 = 0.3 1 = >16	0.23	2	Penicillin G	≤0.015-16	0.125	8
Clindamycin	4-8	8	8	Amoxicillin clavulanate	≤0.015-16	0.15	16
Childaniyeni	<del>-</del> -0	0	0	Cefoxitin	≤0.015-16	0.5	16
Fusobacterium nucleatum (12)				Imipenem	≤0.015-1	0.06	1
Gemifloxacin	0.03-0.25	0.125	0.25	Metronidazole	0.25->32	2	4
Trovafloxacin	0.03-0.5	0.03	0.5	Clindamycin	$\leq 0.015 - 1$	0.25	0.5
Sitafloxacin	≤0.015-0.06	0.03	0.03	Peptostreptococcus asaccharolyticus			
Grepafloxacin	0.25 - 1	0.5	0.5	(13)			
Levofloxacin	0.06 - 1	0.5	0.5	Gemifloxacin	≤0.015-125	0.125	0.125
Sparfloxacin	0.25-2	0.5	1	Trovafloxacin	0.06 - 1	0.5	1
Penicillin G	≤0.015	≤0.015	≤0.015	Sitafloxacin	$\leq 0.015 - 0.5$	0.125	0.25
Amoxicillin clavulanate	$\leq 0.015 - 0.06$	$\leq 0.015$	$\leq 0.015$	Grepafloxacin	0.125-2	0.5	2
Cefoxitin	$\leq 0.015 - 0.25$	$\leq 0.015$	0.06	Levofloxacin	0.25-8	4	8
Imipenem Metronidazole	$\leq 0.015 - 0.03$ $\leq 0.06 - 0.5$	≤0.015 0.125	0.03 0.25	Sparfloxacin	0.06-4	0.25	4
Clindamycin	$\leq 0.06 - 0.5$ $\leq 0.015 - 0.06$	0.125	0.25	Penicillin G	$\leq 0.015 - 0.125$	$\leq 0.015$	0.03
Childaniyelli	-0.013-0.00	0.05	0.00	Amoxicillin clavulanate	$\leq 0.015 - 0.06$	0.03	0.06 0.125
Fusobacterium necrophorum (11)				Cefoxitin Imipenem	0.03–0.125 ≤0.015	$0.06 \le 0.015$	$0.125 \le 0.015$
Gemifloxacin	0.125-0.5	0.25	0.5	Metronidazole	≤0.015 0.125–2	≤0.013 1	≤0.013 2
	0.120-0.0	0.23	0.5		0.123-2	1	L

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### TABLE 1—Continued

Organism (no. of isolates) and agent	MIC			Organism (no. of isolates)	MIC		
	Range	50%	90%	and agent	Range	50%	90%
Clindamycin	≤0.015-128	0.06	32	Clindamycin	≤0.015->126	≤0.015	>128
Peptostreptococcus magnus (14)				Prevotella bivia (10)			
Gemifloxacin	$\leq 0.015 - 1$	0.03	0.03	Gemifloxacin	8-16	8	16
Trovafloxacin	0.06-2	0.06	0.25	Trovafloxacin	1-2	2	2
Sitafloxacin	≤0.015-0.25	$\leq 0.015$	0.06	Sitafloxacin	0.125-0.25	0.25	0.25
Grepafloxacin	0.06-2	0.125	0.5	Grepafloxacin	8-16	16	16
Levofloxacin	0.25-4	0.5	4	Levofloxacin	2–4	2	4
Sparfloxacin	0.06-4	0.125	0.25	Sparfloxacin	4-8	4	8
Penicillin G	≤0.015-0.25	0.06	0.125	Penicillin G	0.06-32	4	16
Amoxicillin clavulanate	0.03-0.25	0.125	0.25	Amoxicillin clavulanate	0.06-2	0.25	2
Cefoxitin	0.125 - 1	0.5	0.5	Cefoxitin	0.25-8	1	8
Imipenem	≤0.015-0.125	0.06	0.06	Imipenem	$\leq 0.015 - 0.06$	0.03	0.06
Metronidazole	0.25 - 2	0.5	1	Metronidazole	0.5-8	2	2
Clindamycin	0.06-2	1	2	Clindamycin	$\leq 0.015 -> 128$	$\leq 0.015$	0.03
Peptostreptococcus micros (12)				Prevotella buccae (10)			
Gemifloxacin	0.06-0.125	0.06	0.06	Gemifloxacin	0.5-4	2	2
Trovafloxacin	0.06-0.125	0.06	0.06	Trovafloxacin	0.125 - 2	0.5	1
Sitafloxacin	≤0.015-0.03	≤0.015	0.03	Sitafloxacin	0.03-0.5	0.03	0.06
Grepafloxacin	1	1	1	Grepafloxacin	1-8	4	8
Levofloxacin	0.5 - 1	0.5	0.5	Levofloxacin	0.5->16	0.5	1
Sparfloxacin	0.5	0.5	0.5	Sparfloxacin	1->16	1	2
Penicillin G	≤0.015-0.06	≤0.015	0.03	Penicillin G	0.06 -> 32	16	>32
Amoxicillin clavulanate	0.06-0.5	0.06	0.25	Amoxicillin clavulanate	0.06-1	0.25	0.5
Cefoxitin	0.5-1	0.5	1	Cefoxitin	0.5–16	4	8
Imipenem	≤0.015-0.03	0.03	0.03	Imipenem	0.03-0.125	0.06	0.125
Metronidazole	≤0.06-0.5	0.05	0.05	Metronidazole	1-8	2	2
Clindamycin	0.125–1	0.125	0.5	Clindamycin	$\leq 0.015 - 0.25$	$\leq 0.015$	≤0.015
Peptostreptococcus prevotii (14)				Prevotella intermedia (10)			
Gemifloxacin	0.03-16	0.06	0.25	Gemifloxacin	0.06-0.5	0.5	0.5
Trovafloxacin	0.126->16	0.00	1	Trovafloxacin	0.03-0.5	0.5	0.5
Sitafloxacin	≤0.015-1	0.25	0.5	Sitafloxacin	≤0.015-0.06	0.5 ≤0.015	0.03
Grepafloxacin	0.25 > 16	0.00	4		≤0.013=0.00 0.5-2	≤0.015 1	2
Levofloxacin	$0.25 \rightarrow 10$ $0.25 \rightarrow 16$	2	16	Grepafloxacin			$0.5^{2}$
Sparfloxacin	0.25 > 10 0.06 - > 16	0.25	4	Levofloxacin	0.25-1	0.25 1	
Penicillin G	$\leq 0.015 - 0.125$	0.23	0.125	Sparfloxacin	0.25-2		1 16
Amoxicillin clavulanate		0.03	0.125	Penicillin G	$\leq 0.015 -> 32$	0.03	
Cefoxitin	$\leq 0.015 - 0.125$	0.03	0.125	Amoxicillin clavulanate	0.03-16	0.06	0.5
	0.03-1			Cefoxitin	0.125-32	0.25	2
Imipenem	$\leq 0.015 - 0.06$	≤0.015	0.06 2	Imipenem	≤0.015-2	0.03	0.06
Metronidazole Clindamycin	$0.25 > 32 \le 0.015 - 128$	0.5 0.5	64	Metronidazole Clindamycin	$0.5-2 \le 0.015-0.5$	$1 \le 0.015$	$2 \le 0.015$
Porphyromonas asaccharolytica				-			
1				Prevotella melaninogenica (11)	0.105.16	1	4
(11) Comiflorencin	0.06-0.125	0 1 25	0.125	Gemifloxacin	0.125-16	1	1
Gemifloxacin		0.125	0.125	Trovafloxacin	0.03-4	1	1
Trovafloxacin Site flows size	0.125-0.5	0.25	0.25	Sitafloxacin	≤0.015-0.5	0.03	0.06
Sitafloxacin	$\leq 0.015 - 0.06$	0.03	0.03	Grepafloxacin	0.125-8	0.25	0.5
Grepafloxacin	0.5-4	2	2	Levofloxacin	0.25-8	1	1
Levofloxacin	0.06-1	0.5	0.5	Sparfloxacin	0.25-2	2	2
Sparfloxacin	0.06-1	1	1	Penicillin G	0.03-16	4	8
Penicillin G	≤0.015-2	≤0.015	1	Amoxicillin clavulanate	≤0.015-2	0.125	2
Amoxicillin clavulanate	≤0.015-0.125	≤0.015	0.125	Cefoxitin	0.06-2	1	2
Cefoxitin	$\leq 0.015 - 2$	0.125	1	Imipenem	$\leq 0.015 - 0.06$	0.03	0.06
Imipenem	≤0.015	≤0.015	≤0.015	Metronidazole	0.25-2	0.5	1
Metronidazole	$\leq 0.06 - 0.5$	0.25	0.5	Clindamycin	$\leq 0.015 -> 128$	≤0.015	0.03

<sup>a</sup> Includes F. mortiferum (three strains), F. necrogenes (four strains), and F. ulcerans (five strains).

Cormicon and Jones (1), using brucella blood agar, tested a limited number of *B. fragilis* group species strains and noted that the MIC of gemifloxacin at which 90% of the isolates were inhibited ( $MIC_{90}$ ) was 8 µg/ml but did not identify differences among the various species. Our study found marked differences in the activity of gemifloxacin against the various species

belonging to the *B. fragilis* group. Gemifloxacin was active against *B. fragilis* at  $\leq 2 \mu g/ml$  and was more active than grepafloxacin and levofloxacin (MIC<sub>90</sub>s, 4  $\mu g/ml$ ). Similar results were reported by Marco et al. (4), who also used brucella agar supplemented with 5% sheep blood and an inoculum of 10<sup>5</sup> CFU/ml. They tested 35 strains of *B. fragilis* and reported a MIC range of 0.5 to 8 µg/ml and a MIC<sub>90</sub> of 1 µg/ml for gemifloxacin and a MIC<sub>90</sub> of 0.5 µg/ml (range,  $\leq 0.25$  to 4 µg/ml) for trovafloxacin. They did not, however, report results for other individual member species of the *B. fragilis* group. In our study, sitafloxacin was generally four times more active (MIC<sub>90</sub>, 0.25 µg/ml) than gemifloxacin, and trovafloxacin (MIC range, 0.125 to 4 µg/ml; MIC<sub>90</sub>, 0.5 µg/ml) was twice as active. For almost all *Bacteroides distasonis* and *Bacteroides ovatus* strains, the MIC of gemifloxacin was >2 µg/ml. For all but one *Bacteroides stercoris* strain, the MIC of gemifloxacin was  $\leq 0.5$  µg/ml. *B. thetaiotaomicron* and *Bacteroides caccae* strains had a biphasic distribution of susceptibility to gemifloxacin, which had a MIC<sub>50</sub> of 1 µg/ml but a MIC<sub>90</sub> of 16 µg/ml. *Bacteroides uniformis* and *Bacteroides vulgatus* showed marked strain variation in relation to gemifloxacin.

All *Prevotella intermedia* strains and all but one strain of *Prevotella melaninogenica* were susceptible to  $\leq 0.5$  and  $\leq 1 \mu g$  of gemifloxacin/ml, respectively. Generally,  $\geq 2 \mu g/ml$  was required for inhibition of *Prevotella buccae* and  $\geq 8 \mu g/ml$  for that of *Prevotella bivia*. The only species of *Porphyromonas* that we tested was *Porphyromonas asaccharolytica*, all strains of which were inhibited by  $\leq 0.125 \mu g$  of gemifloxacin/ml.

Gemifloxacin showed generally good activity against grampositive anaerobic bacteria. The respective MIC<sub>90</sub>s of gemifloxacin, trovafloxacin, and sitafloxacin against the various clostridia were as follows: for Clostridium clostridioforme, 0.5, 8, and 0.25  $\mu$ g/ml; for *Clostridium difficile*, >16, >16, and 1  $\mu$ g/ ml; for Clostridium innocuum, 2, 4, and 1 µg/ml; for Clostridium perfringens, 0.06, 0.25, and 0.06 µg/ml; and for Clostridium ramosum, 8, 8, and 4 µg/ml. Marco et al. (4) noted that gemifloxacin and trovafloxacin had MIC<sub>90</sub>s of 1 µg/ml against a mélange of clostridial species. They also noted that sparfloxacin was much less active, with a  $MIC_{90}$  of 8 µg/ml. In our study, gemifloxacin was 1 to 2 dilutions more active than trovafloxacin against fusobacteria and peptostreptococci (Peptostreptococcus anaerobius, Peptostreptococcus asaccharolyticus, Peptostreptococcus magnus, Peptostreptococcus micros, and Peptostreptococcus prevotii) and was equivalent to trovafloxacin in activity against P. asaccharolytica and clostridia. Gemifloxacin was equivalent to sitafloxacin against peptostreptococci, C. perfringens, and C. ramosum and was 2 to 3 dilutions less active against fusobacteria. Sparfloxacin, grepafloxacin, and levofloxacin were generally less active than gemifloxacin against all anaerobes. Marco et al. (4) reported that gemifloxacin and trovafloxacin each had a MIC<sub>90</sub> of 2 µg/ml against a combined group of 18 strains of peptostreptococci and a MIC<sub>90</sub> of 4  $\mu$ g/ml against fusobacteria. Because the study by Marco et al. (4) used 119 strains of anaerobes of various genera but did not report data on individual species other than *B. fragilis*, it is difficult to draw comparisons with our results, which found such marked variation among species.

Gemifloxacin had selectively potent activity against the various anaerobic species tested, with generally improved activity against gram-positive anaerobes and fusobacteria. It showed moderate but variable activity against gram-negative anaerobes. The advantage of this selective anaerobic activity may become evident if gemifloxacin is proven to have clinical efficacy in situations such as dental, head and neck, and pleuropulmonary infections, where gram-positive anaerobes, fusobacteria, and some *Prevotella* and *Porphyromonas* species predominate, while minimizing disturbance of the normal enteric flora.

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