Antimicrobial Activity of Garenoxacin (GRN), a Novel Des-F(6) Quinolone Tested Against 29,460 Community-Acquired Respiratory Pathogens (CA-RTI) from Europe, North America and Latin America



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Abstract

Background: Contemporary (1999-2005) activity of GRN and comparison fluoroquinolones (FQs) were tested against S. pneumoniae (SPN), H. influenzae (HI) and M. catarrhalis (MCAT) isolates from 3 continents. Trends in resistance and effects of co-R on GRN potency was assessed.

Methods: SPN (13,273), HI (11,695) and MCAT (4,492) were collected from more than 100 laboratories worldwide and tested by CLSI methods in a central reference laboratory, with QC results within published CLSI ranges. Subsets with penicillin-R (PEN; ≥2 mg/L, SPN), levofloxacin-R (LEVO; ≥4 mg/L, SPN) and ampicillin-R (AMP; ≥2 mg/L, HI) were analyzed separately.

Results: GRN exhibited excellent potency (MIC₉₀, ≤0.06 mg/L) against all 3 CA-RTI associated species, not differing by continent or Co-R patterns among PEN or AMP-R subsets. GRN MIC results by continent are shown in the table.

Organism	Continent (no) ^a	MIC_{90}	Highest MIC	% ≤1 mg/L
SPN	NA (6909)	0.06	>4	99.86
	LA (1554)	0.06	2	99.94
	EU (4810)	0.06	2	99.96
HI	NA(6069)	≤0.03	16	99.97
	LA (1178)	≤0.03	0.12	100.0
	EU (4448)	≤0.03	0.25	100.0
MCAT	NA (2685)	≤0.03	0.12	100.0
	LA (289)	≤0.03	0.12	100.0
	EU (1518)	≤0.03	0.12	100.0

a. NA, N. America; LA, Latin America; EU, Europe

All GRN MICs for MCAT were ≤0.03 mg/L and only one HI strain was FQ-R (GRN S rate at 99.99%). 13 SPN strains had GRN MICs at >1 mg/L (0.09%) with only 3 strains with a MIC of \geq 4 mg/L (0.02% of SPN). No trend toward greater GRN-R was noted over time (1999-2005), and GRN was 4- and 16-fold more active than moxifloxacin and LEVO, respectively, versus SPN. LEVO-R rate among SPN was 0.95%, 10-fold greater than GRN. GRN MICs (MIC₉₀, 1 mg/L) for LEVO-R SPN were 16-fold higher, making the MIC₉₀ for GRN against LEVO-R SPN the same as the MIC₉₀ for LEVO against the wild-type (non-mutant) population of SPN.

Conclusions: GRN is a very potent investigational agent with an activity and spectrum at ≤1 mg/L, highest among tested FQ agents. Rare GRN-R SPN and HI strains (≤0.1%) were noted, each having ≥4 QRDR mutations (data not shown). GRN should be a welcome advance for CA-RTI therapy.

Introduction

Garenoxacin is a novel des-F(6)-quinolone that lacks the C6position fluorine and has a unique difluoromethoxy substitution at position C8. These alterations resulted in a quinolone with improved potency against both DNA gyrase and topoisomerase IV. Garenoxacin has been described as highly active against important Grampositive and -negative pathogens including: Enterobacteriaceae, staphylococci, streptococci (S. pneumoniae, viridans group species and B-haemolytic streptococci), Acinetobacter spp. and some other Gram-negative non-fermentative bacilli, Haemophilus influenzae, Moraxella catarrhalis, atypical respiratory tract pathogens (mycoplasmas, C. pneumoniae, and Legionella spp.), many enterococci and anaerobes, especially Gram-positive species. These features are complemented by a favorable pharmacokinetic/pharmacodynamic profile, leading to a high AUC/MIC ratio. This high AUC/MIC ratio predicts a greater probability of favorable target attainment that has been associated with successful bacterial eradication and minimization of mutational events among indicated species (low MPC values). These elements of spectrum and potency favor garenoxacin applications for community-acquired respiratory tract infections (CA-RTI; hospitalized or ambulatory patients).

Since garenoxacin features high potency and breadth of spectrum against common CA-RTI pathogens (29,460 strains tested since 1999 on three continents [Europe, North and Latin America]), subsets of strains having elevated MIC values to currently utilized agents were selected for further studies and compared to all studied strains of *S. pneumoniae*, *H. influenzae*, and *M. catarrhalis*. Garenoxacin MICs versus those organisms and the determination of QRDR mutations for some isolates were used to establish the role of this investigational des-F(6)-quinolone against emerging resistant pathogen types.

Materials and Methods

Bacterial strains tested: Recent RTI strains were categorized into those treated in the ambulatory setting (CA-RTI). The organisms were cultured during 1999-2005 from Latin America (10 medical centers), North America (USA and Canada; 23) and Europe (20). The number of strains was: *S. pneumoniae* (13,273), *H. influenzae* (11,695) and M. catarrhalis (4,492). These CA-RTI isolates came from non-duplicated cultures in 23 nations.

Susceptibility testing methods: All MIC values were generated using broth microdilution methods (CLSI M7-A7) with panels produced by TREK Diagnostics (Cleveland, Ohio, USA). Cationadjusted Mueller-Hinton broth was supplemented where indicated with 2-5% lysed horse blood (streptococci) and HTM components (Haemophilus). Concurrent quality assurance was maintained via use of CLSI-recommended strains including Escherichia coli ATCC 25922 and 35218, Pseudomonas aeruginosa ATCC 27853, Enterococcus faecalis ATCC 29212, Staphylococcus aureus ATCC 25923 and 29213, *H. influenzae* ATCC 49247 and 49766, and

S. pneumoniae ATCC 49619. All quality control results were within published MIC ranges (M100-S16) for each agent tested. More than 30 different antimicrobial agents were processed each year with selected agents compared to garenoxacin in this presentation (see Tables). A breakpoint for garenoxacin susceptibility and resistance at $\leq 1/\geq 4$ mg/L was used for comparison purposes only.

Molecular methods: The QRDR was assessed for mutations in gyrA or gyrB and, parC or parE by PCR amplication and sequence analyses. S. pneumoniae and H. influenzae isolates with elevated garenoxacin MIC results (2->4 mg/L) were processed to detect mutations as were selected isolates with fluoroquinolone MICs ≥ three log₂ dilutions above the wild-type distributions. A total of 124 isolates had QRDR sequences determined.

Results

- A large amount of quantitative (MIC values) activity information has been accumulated on garenoxacin against CA-RTI pathogens (29,460 strains; Tables 1-3) from three continents.
- No significant variations in garenoxacin potency were observed among S. pneumoniae (MIC_{50 and 90}, 0.06 mg/L), *H. influenzae* (MIC₉₀, \leq 0.03 mg/L) and *M. catarrhalis* (MIC₉₀, \leq 0.03 mg/L) between strains originating from Europe, North America or Latin America.
- Only two *H. influenzae* strains (North America), and 12 *S. pneumoniae* strains (two in Europe, nine in North America, one in Latin America) were detected with a garenoxacin MIC at >1 mg/L. All potentially garenoxacin-nonsusceptible strains had multiple [≥3] mutations in the QRDR (data not shown).

Activity of garenoxacin tested against 29,460 CA-Table 1. RTI pathogens and listed by continent or region of isolation (SENTRY Program, 1999-2005).

Organism (No. tested/	Eur	rope	North A	America	Latin /	America
Antimicrobial agent)	MIC ₅₀ /MIC ₉₀	% susceptible	MIC ₅₀ /MIC ₉₀	% susceptible	MIC ₅₀ /MIC ₉₀	% susceptibl
H. influenzae	(4,4	148)		069)	(1,	178)
Garenoxacin	≤0.03/≤0.03	100.0	≤0.03/≤0.03	>99.9	≤0.03/≤0.03	100.0
Ciprofloxacin	≤0.03/≤0.03	100.0	≤0.03/≤0.03		≤0.03/≤0.03	100.0
Levofloxacin	≤0.03/≤0.03	100.0	≤0.03/≤0.03	>99.9	≤0.03/≤0.03	100.0
Moxifloxacin	≤0.03/≤0.03	100.0	≤0.03/≤0.03	>99.9	≤0.03/≤0.03	100.0
Ampicillin	≤0.5/>4	84.0	≤0.5/>4	72.1	≤0.5/>4	84.5
Amox/clav ^b	0.5/1	99.8	0.5/1	>99.9	0.5/1	100.0
Azithromycin	1/2	99.8	1/2	99.6	1/2	99.8
Cefuroxime axetil	1/2	98.2	1/2	98.7	1/2	98.7
Tetracycline	≤2/≤2	97.3	≤2/≤2	99.4	≤2/≤2	96.5
Trim/sulfab	≤0.5/>4	79.5	≤0.5/>4	79.3	≤0.5>4	61.1
M. catarrhalis	(1,!	518)	(2,6	585)	(2	89)
Garenoxacin	≤0.03/≤0.03	_C	≤0.03/≤0.03	-	≤0.03/≤0.03	
Ciprofloxacin	≤0.03/0.06	-	≤0.03/0.06	-	≤0.03/0.06	-
Levofloxacin	≤0.03/0.06	-	≤0.03/0.06	-	≤0.03/0.06	-
Moxifloxacin	0.06/0.06	-	0.06/0.06	-	0.06/0.06	-
Penicillin	4/>4	(4.9) ^d	>4/>4	(4.8)	>4/>4	(2.8)
Amox/clav ^b	≤0.25/≤0.25	-	≤0.25/≤0.25		≤0.25/≤0.25	-
Erythromycin	≤0.25/≤0.25	_	≤0.25/≤0.25		≤0.25/≤0.25	_
Cefuroxime axetil	1/2	_	1/2	-	1/2	_
Tetracycline	≤2/≤2	_	≤2/≤2	-	≤2/≤2	
Trim/sulfa ^b	≤0.5/≤0.5	-	≤0.5/≤0.5	-	≤0.5/≤0.5	-
S. pneumoniae	(4.8	310)	(6.9	909)	(1,554)	
Garenoxacin	0.06/0.06	>99.9	0.06/0.06	>99.9	0.06/0.06	>99.9
Ciprofloxacin	1/2	(4.6)e	1/2	(3.2)	1/2	(3.3)
Levofloxacin	1/1	99.0	1/1	98.9	1/1	99.7
Moxifloxacin	0.12/0.25		0.12/0.25		0.12/0.25	
Penicillin	≤0.03/2		≤0.03/2		≤0.03/2	70.9
Cefuroxime axetil	≤1/4		≤1/4	75.4	≤1/4	85.1
Ceftriaxone	≤0.25/1	97.6	≤0.25/1		≤0.25/1	
Amox/clav ^b	≤1/2		≤1/2		≤1/2	97.4
Erythromycin	≤0.25/>2		≤0.25/>2		≤0.25/>2	
Clindamycin	≤0.25/>2		≤0.25/≤0.25		≤0.25/≤0.25	
Tetracycline	≤2/>8		≤2/>8		≤2/>8	81.6
Trim/sulfab	≤0.5/>2	65.8	≤0.5/>2		≤0.5/>2	55.0

B-lactamase-positive test results were used to determine resistance e. Percentage in parentheses indicate the ciprofloxacin MICs at ≥4 mg/L, possible QRDR mutants.

Impact of various resistances in CA-RTI pathogens Table 2. on the activity of garenoxacin in three geographic regions.

				renoxacin MICs (mg/L) by region					
Organism/resistant	Europe			North America			Latin America		
group (no. tested)	50%	90%	Range	50%	90%	Range	50%	90%	Range
H. influenzae	(3,728/710)			(4,368/1,690)			(995/183)		
Ampicillin-susceptible	≤0.03	≤0.03	≤0.03-0.25	≤0.03	≤0.03	≤0.03-16 ^a	≤0.03	≤0.03	≤0.03-0.12
Ampicillin-resistant	≤0.03	≤0.03	≤0.03-0.12	≤0.03	≤0.03	≤0.03-0.12	≤0.03	≤0.03	≤0.03-0.12
M. catarrhalis	arrhalis (69/1,449)		(125/2,560)			(7/282)			
ß-lactamase-negative	≤0.03	≤0.03	≤0.03-0.12	≤0.03	≤0.03	≤0.03	≤0.03	_	≤0.03
ß-lactamase-positive	≤0.03	≤0.03	≤0.03-0.06	≤0.03	≤0.03	≤0.03-0.12	≤0.03	≤0.03	≤0.03-0.12
S. pneumoniae	(3	339/727	/750)	(4,2	237/1,16	53/1,225)	(1,004/2	58/191)
Penicillin-susceptible	0.06	0.06	≤0.03-2 ^b	0.06	0.06	≤0.03-4°	0.06	0.06	≤0.03-2 ^d
Penicillin-intermediate	0.06	0.06	≤0.03-1	0.06	0.06	≤0.03-2 ^e	0.06	0.06	≤0.03-0.12
Penicillin-resistant	0.06	0.06	≤0.03-1	0.06	0.06	≤0.03->4 ^f	0.06	0.06	≤0.03-0.12
a. Two strains with garen b. Two strains at 2 mg/L. c. One strain each at 2 add. One strain at 2 mg/L. e. Four strains at 2 mg/L	nd 4 mg		lts at 2 and 16	5 mg/L, n	nultiple	QRDR mutatio	ns confir	med.	

f. One strain at 2 and two isolates at >4 mg/L.

Among 126 levofloxacin-nonsusceptible *S. pneumoniae* (MIC, ≥4 mg/L), garenoxacin MIC values were ≤1 mg/L for nearly 90% of strains (MIC₅₀, 0.5 mg/L).

MIC distributions of five quinolones tested against Table 3. 126 levofloxacin non-susceptible (MIC, ≥4 mg/L) S. pneumoniae strains from the SENTRY Program (1999-2005).

Cum. % inhibited at MIC (mg/L)									
Fluoroquinolone	≤0.03	0.06	0.12	0.25	0.5	1	2	4	% Susceptible ^a
Garenoxacin	2.4	4.8	9.5	18.3	55.6	89.7	97.6	98.4	89.7 ^b
Ciprofloxacin	0.0	0.0	0.0	0.0	0.0	0.0	0.8	-C	_d
Gatifloxacin	0.0	0.0	0.0	0.0	0.8	3.2	14.3	78.6	3.2
Levofloxacin	0.0	0.0	0.0	0.0	0.0	0.0	0.0	9.5	0.0
Moxifloxacin	0.0	0.0	0.0	2.7	4.5	14.4	67.6	94.6	14.4

a. CLSI (2006) criteria for susceptibility. b. Susceptible percentage for a breakpoint of only ≤ 1 mg/L. c. - = untested concentration.

d. - = no criteria for this species (CLSI, 2006).

Conclusions

- Garenoxacin in vitro susceptibility testing results for nearly 30,000 CA-RTI pathogens revealed a potential resistance rate of only 0.05% (MIC, > 1 mg/L).
- Garenoxacin should be a welcome addition for CA-RTI therapy, especially for use against emerging isolates that are resistant to Blactams, MLS_B class agents, trimethoprim/sulfamethoxazole, tetracyclines, and recently introduced fluoroquinolones, such as levofloxacin.

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