

Pharmacodynamic evaluation of the Intracellular activity of CEM-101, a novel fluoroketolide, towards *Staphylococcus aureus*, *Listeria monocytogenes*, and *Legionella pneumophila* in human THP-1 macrophages.

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Abstract (revised)

Background:

Macrolides accumulate in eukaryotic cells, which is considered advantageous for the treatment of intracellular infections. Ketolides, i.e. macrolides in which the cladinose has been removed and replace by a ketone group, recover activity against most macrolide-resistant organisms. CEM-101 is a novel fluoroketolide with a 11,12 carbamate-butyl-[1,2,3]-triazolyl-phenylamino side chain, which demonstrates enhanced potency compared to telithromycin. We have assessed the cellular accumulation and intracellular activity of CEM-101 against the intracellular forms of *Staphylococcus aureus* (*S.a.*), *Listeria monocytogenes* (*L.m.*), and *Legionella pneumophila* (*L.p.*) in comparison with azithromycin (AZM), clarithromycin (CLR), and telithromycin (TEL).

Methods:

All experiments were performed with the human macrophage cell line THP-1. Drug accumulation was measured using a bioassay. Intracellular activity was measured over time and concentration by following the change in cell-associated CFU compared to post-phagocytosis levels (see details in JAC 2004;54:288-9 [L.m. - strain EGD]; AAC 2006; 50:841-51 [S.a. strain ATCC 25923]; similar protocol for *L.p.* [strain ATCC 33153]).

Results:

Uptake of CEM-101 was linear over time, reaching accumulation levels about 375-fold within 24 h (AZM, 160 μ M, 30 μ M, TEL, 21 μ M). Accumulation was suppressed by acid pH or addition of the proton ionophore monensin, but not modified by verapamil or gemfibrozil (specific inhibitors of P-gp and MRP, respectively). MIC and intracellular activities (developed in a concentration-dependent fashion [Hill equation] on the Table).

CEM-101		AZM		CLR		TEL						
MIC *	Cs ^b	E _{max} *	MIC *	Cs ^b	E _{max} *	MIC *	Cs ^b	E _{max} *				
S.a.	0.06	0.022	-0.86	0.5	> 20	0.04	0.5	0.84	-0.18	0.25	0.63	-0.29
L. m.	0.004	0.11	-0.66	1	11.6	-0.81						
L. p.	0.004	0.008	-1.03	0.016	2.90	-0.83	0.007	0.12	-0.71	0.007	0.06	-0.63

* mg/L; ^b static concentration (mg/L) at 24 h; ^c log₁₀ CFU at 24 h compared to the post-phagocytosis inoculum

CEM-101 showed similar or slightly higher relative maximal efficacy (E_{max}) compared to AZM, CLR or TEL, but considerably higher relative potency (lower E_{max} and C_s), in its lower MIC when expressed on a mass basis (differences in E_{max} and C_s between drugs largely vanish if data are expressed as multiples of the MIC).

Conclusions: CEM-101 is a fluoroketolide with enhanced cellular accumulation. It shows improved intracellular potency (on a weight basis) in comparison with AZM, CLR and TEL in this in vitro model (mainly due to its larger intrinsic activity [lower MICs] against target organisms). This should lead to enhanced in vivo potency if used doses similar to those of the comparators tested here.

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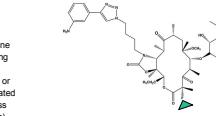
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Introduction

Macrolides accumulate inside eukaryotic cells, which has been considered advantageous for the treatment of intracellular infections (1,2), even though we know that these antibiotics express only a minimal fraction of their antibacterial potential intracellularly (3).

11,12-carbamate analogs of clarithromycin carrying a lipophilic side chain show improved activity compared to the parent compound (4). Together with the removal of the cladinose, this led to the discovery and development of the telithromycin (5), the first ketolide to reach clinical approval. CEM-101 is a novel fluoroketolide containing an 11,12-carbamate-butyl-[1,2,3]-triazolyl-aminophenyl side chain (see structure hereunder) that shows enhanced potency compared to telithromycin (6). We have assessed its cellular accumulation and intracellular activity using models that have been developed for the study of the intracellular pharmacodynamics of antibiotics (3,7).

Structural formula of CEM-101



The molecule possesses a 11,12-carbamate-butyl-[1,2,3]-triazolyl-aminophenyl sidechain and a fluorine atom (green arrowhead) substituting the 11-position of the core molecule. It is monocationic at neutral or moderately acidic pH, as the calculated pKa of the aminophenyl group is less than 4 (vs. ~8.5 for the desosamine).

Methods

Cells: THP-1 myelomonocytic cells (ATCC TIB-202) with macrophage-like activity (7).

Antibiotic assay: microbiological method (7) and accumulation calculated as apparent intracellular/extracellular concentration ratio (7) based on cell protein content (7)

Bacterial strains: *S. aureus* ATCC 25923 and *L. monocytogenes* strain EGD and *L. pneumophila* strain ATCC 33153

Infection and assessment of antibiotic intracellular activities: Time- (0-24h) and concentration-dependent (25) *S. aureus* and *L. monocytogenes*, 48h for *L. pneumophila* experiments described in ref. 3. Preparation of THP-1 macrophages. Data from concentration-dependent experiments using Hill's equation (7) to calculate pharmacological descriptors (E_{max}: maximal reduction of the intracellular inoculum (log₁₀ units) for an infinitely large antibiotic concentration; E_{min}: increase in intracellular inoculum (log₁₀ units) for an infinitely low antibiotic concentration; EC₅₀: antibiotic concentration yielding a response halfway between E_{min} and E_{max}; C_s: antibiotic concentration yielding a static effect).

Results

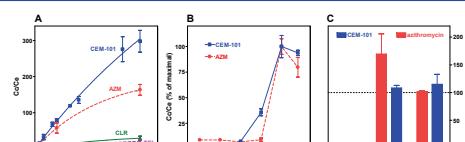


Figure 1: Accumulation of CEM-101 vs. comparators in uninfected cells at 37°C
(*C = apparent cellular concentration [in mg/L], based on cell protein assay; Cs = extracellular concentration [10 mg/L]).
A. Influence of incubation time; B. Influence of pH (30 min); C. Influence of monensin (10 μ M, 2 h) incubation; verapamil (P-gp inhibitor, 100 μ M, 24 h); gemfibrozil (MRP inhibitor, 250 μ M, 24 h)

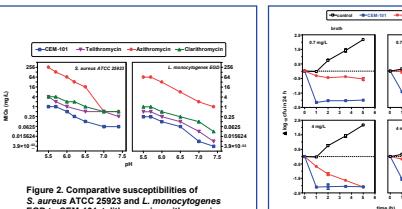


Figure 2: Comparative susceptibilities of *S. aureus* ATCC 25923 and *L. monocytogenes* EGD to CEM-101, telithromycin, azithromycin, and clarithromycin.
MIC determinations in pH-adjusted broth.

MIC determinations in pH-adjusted broth.

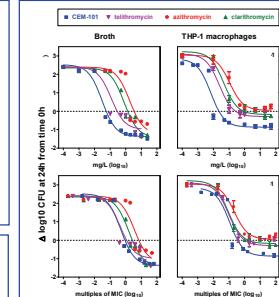


Figure 3: Short-term kill effect of CEM-101 and azithromycin towards *S. aureus* (ATCC 25923) in broth vs. in THP-1 macrophages.
Change in CFU (A log₁₀ CFU at 24 h / CFU at 0 h) per mg of cell protein at 24 h (THP-1 macrophages) or 48 h (*L. pneumophila*) compared to initial post-phagocytosis inoculum.
Abscissae: weight concentrations (upper rows) or multiples of MIC (measured at pH 7.4; lower rows).

Figure 4: Concentration-effect relationships for CEM-101, telithromycin, clarithromycin, and azithromycin against *S. aureus* (ATCC 25923) in broth or in THP-1 macrophages.
Ordinate: change in CFU (A log₁₀ CFU) per mL (broth) or per mg of cell protein (THP-1 macrophages) at 24 h compared to initial inoculum;
Abscissae: weight concentrations (upper rows) or multiples of MIC (measured at pH 7.4; lower rows).

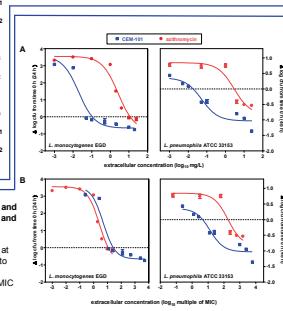


Figure 5: Concentration-effect relationships for CEM-101 and azithromycin towards intraphagocytic *L. monocytogenes* and *L. pneumophila*.
Change in CFU (A log₁₀ CFU at 24 h / CFU at 0 h) per mg of cell protein at 24 h (*L. monocytogenes* EGD) or 48 h (*L. pneumophila*) compared to initial post-phagocytosis inoculum.
Abscissae: weight concentrations (upper rows) or multiples of MIC (measured at pH 7.4; lower rows).

(data from Figure 4: E_{max} = A log₁₀ CFU compared to post-phagocytosis inoculum; EC₅₀ and C_s; mg/L or x MIC)

antibiotic	broth			THP-1 macrophages		
	E _{max}	EC ₅₀	C _s	E _{max}	EC ₅₀	C _s
CEM-101	-1.37	mg/L 0.03	0.06	-0.86	mg/L 0.0068	0.022
telithromycin	-1.00	mg/L 0.12	0.29	-0.29	mg/L 0.024	0.63
azithromycin	-1.23	mg/L 1.78	3.4	0.04	mg/L 0.11	> 50
clarithromycin	-1.41	mg/L 0.80	1.32	-0.18	mg/L 0.046	0.84
		x MIC 3.95			x MIC 0.22	> 100
		x MIC 1.59	2.65		x MIC 0.093	1.68

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