

# CEM-101, a Novel Ketolidide, *In Vitro* Activity Against Resistant Strains of *Streptococcus pneumoniae* and *Haemophilus influenzae*

J. DUBOIS<sup>1\*</sup>, P. FERNANDES<sup>2</sup><sup>1</sup>M360, Sherbrooke, Canada, <sup>2</sup>Cempra Pharmaceuticals Inc., Chapel Hill, USA30 th ECCMID, Vienna, Austria  
Jacques Dubois Ph.D.  
M360, Sherbrooke, Québec, Canada  
819.571.4366 fax 819.843.1391  
jdubois@m360.ca

## Abstract

**Objective:** CEM-101 is a promising fluoroketolidide that has potent activity against respiratory tract pathogens resistant to other macrolide agents. Its activity against a variety of resistant strains of *Streptococcus pneumoniae* and *Haemophilus influenzae* was investigated.

**Methods:** The *in vitro* activity of CEM-101 was compared with that of telithromycin, azithromycin, erythromycin, levofloxacin and doxycycline against a total of 199 resistant *S. pneumoniae* and 191 resistant *H. influenzae* by agar dilution procedures (CLSI, M7-A7, M100-S18). The tested strains included *S. pneumoniae* erythromycin-resistant (*erm* B genotype; 107 isolates and *mefE* genotype; 54) and ciprofloxacin-resistant (*gyrA* and *parC* genotype; 38) and also *H. influenzae* erythromycin-resistant (*erm* A,B,C genotype; 138) and cipro-resistant (*gyrA* and *parC* genotype; 53).

**Results:** Against *S. pneumoniae* ery-resistant strains (*ermB* genotype), the activity of CEM-101 (MIC<sub>90</sub> 1mg/L) and levofloxacin (MIC<sub>90</sub> 2mg/L) was superior to the macrolides tested: telithromycin (MIC<sub>90</sub> 4mg/L), azithromycin (MIC<sub>90</sub> ≥64 mg/L), erythromycin (MIC<sub>90</sub> ≥64 mg/L) and doxycycline (MIC<sub>90</sub> 32 mg/L). Against *S. pneumoniae* ery-resistant (*mefE* genotype) group, CEM-101 (MIC<sub>90</sub> 0.25 mg/L) was the most active agent followed by levofloxacin (MIC<sub>90</sub> 2mg/L), telithromycin (MIC<sub>90</sub> 8 mg/L), doxycycline (MIC<sub>90</sub> 16 mg/L), azithromycin (MIC<sub>90</sub> ≥64 mg/L) and erythromycin (MIC<sub>90</sub> ≥64 mg/L). Against *S. pneumoniae* cipro-resistant (*gyrA* and *parC* genotype) group, CEM-101 (MIC<sub>90</sub> 0.25 mg/L) was also the most active agent tested followed by telithromycin (MIC<sub>90</sub> 1 mg/L), levofloxacin (MIC<sub>90</sub> 2mg/L), doxycycline (MIC<sub>90</sub> 16 mg/L), azithromycin (MIC<sub>90</sub> ≥64 mg/L) and erythromycin (MIC<sub>90</sub> ≥64 mg/L). Against *H. influenzae* ery-resistant (*erm*A,B,C genotype) strains, CEM-101 (MIC<sub>90</sub> 4 mg/L) was the most active macrolide tested followed by telithromycin (MIC<sub>90</sub> 16 mg/L), azithromycin (MIC<sub>90</sub> 16 mg/L) and erythromycin (MIC<sub>90</sub> ≥64 mg/L). Against *H. influenzae* cipro-resistant (*gyrA* and *parC* genotype) group, CEM-101 (MIC<sub>90</sub> 2 mg/L) was slightly more active than telithromycin (MIC<sub>90</sub> 4 mg/L) and levofloxacin (MIC<sub>90</sub> 4 mg/L).

**Conclusions:** These data confirm the interesting activity of the new fluoroketolidide CEM-101 against resistant *Streptococcus pneumoniae* and *Haemophilus influenzae*.

## Introduction

CEM-101 is a novel fluoroketolidide antibacterial agent related to 14-membered ring macrolides. CEM-101 appears to exhibit superior ability to bind to the ribosomes dimethylated at A2058 by the action of *erm* methyltransferase.

In susceptibility studies, CEM-101 is appreciably more potent than most macrolides or azalides against many Gram-positive organisms, including resistant *Streptococcus pneumoniae*, *Streptococcus pyogenes* and *Staphylococcus* spp. It has potent activity against various atypical respiratory pathogens like *Legionella pneumophila*, *Mycoplasma* spp. and *Chlamydia* spp.

## Objective

We determined the minimum inhibitory concentration (MIC) of CEM-101, telithromycin, azithromycin, erythromycin, levofloxacin and doxycycline against a variety of *Streptococcus pneumoniae* and *Haemophilus influenzae* strains isolated from patient sources.

## Materials and Methods

### Strains

- A variety of recent strains (1995-2008) of *Streptococcus pneumoniae* and *Haemophilus influenzae* were isolated, mostly from upper or lower respiratory tract or blood culture.
- Multiple cultures from the same patient or source were excluded unless a change in organism or antibiogram was noted.
- Organisms were identified by standard methods such as described by Murray et al (1).

Microorganisms	Number of tested strains
<i>Streptococcus pneumoniae</i>	199
-Erythromycin-resistant ( <i>mefE</i> genotype)	107
-Erythromycin-resistant ( <i>erm</i> B genotype)	54
-Ciprofloxacin-resistant ( <i>gyrA</i> and <i>parC</i> genotype)	38
<i>Haemophilus influenzae</i>	191
-Erythromycin-resistant ( <i>erm</i> A, B, C genotype)	138
-Ciprofloxacin-resistant ( <i>gyrA</i> and <i>parC</i> genotype)	53

### Determination of MICs

- MICs were determined using the CLSI agar dilution method (2, 3), with replicate plating of the organisms onto a series of agar plates of increasing concentrations from 0.004 mg/L to 64 mg/L.
  - Mueller-Hinton agar was used as the medium against *S. aureus* strains.
  - Staphylococcus aureus* ATCC25923 and *Escherichia coli* ATCC25922 were included as controls.
- Determinations of genotype *mecA*, *ermA*, *B*, *C*, *mefE* and *gyrA* and *parC***
- Genomic DNA was isolated as described by Smith *et al* (4)
  - Multiplex PCR was performed with primers specific for *mecA*, *ermA*, *ermB*, *ermC* and *mefE* as described by Sutcliffe *et al* (5)
  - Multiplex PCR was performed with primers specific for *gyrA* and *parC* as described by Gonzalez *et al* (6)

## Results

TABLE 1. Susceptibility of *Streptococcus pneumoniae*

Organism (no. tested)	Antibiotic	MIC (mg/L)		
		Range	50%	90%
<i>S. pneumoniae</i> Erythromycin-R <i>mefE</i> (107)	CEM-101	0.016-2	0.25	1
	Telithromycin	0.06-32	1	4
	Azithromycin	4-≥64	≥64	≥64
	Erythromycin	0.06-≥64	≥64	≥64
	Levofloxacin	0.25-2	1	2
	Doxycycline	0.06-32	16	32
<i>S. pneumoniae</i> Erythromycin-R <i>erm</i> B (54)	CEM-101	0.008-2	0.06	0.25
	Telithromycin	0.12-8	0.25	8
	Azithromycin	0.008-≥64	4	≥64
	Erythromycin	0.06-≥64	16	≥64
	Levofloxacin	0.5-2	1	2
	Doxycycline	0.12-32	4	16
<i>S. pneumoniae</i> Ciprofloxacin-R <i>gyrA</i> , <i>parC</i> (38)	CEM-101	0.016-0.25	0.03	0.25
	Telithromycin	0.06-2	0.12	1
	Azithromycin	0.12-≥64	0.25	≥64
	Erythromycin	0.12-≥64	0.25	≥64
	Levofloxacin	1-4	2	2
	Doxycycline	0.06-32	0.5	16

## Results continued

TABLE 2. Susceptibility of *Haemophilus influenzae*

Organism (no. tested)	Antibiotic	MIC (mg/L)		
		Range	50%	90%
<i>H. influenzae</i> Erythromycin-R <i>erm</i> A,B,C (138)	CEM-101	0.12-8	4	4
	Telithromycin	0.25-≥64	8	16
	Azithromycin	0.12-≥64	8	16
	Erythromycin	0.25-≥64	32	≥64
	Levofloxacin	0.008-0.016	0.016	0.016
	Doxycycline	0.12-2	0.5	0.5
<i>H. influenzae</i> Ciprofloxacin-R <i>gyrA</i> , <i>parC</i> (53)	CEM-101	0.12-4	1	2
	Telithromycin	0.25-16	2	4
	Azithromycin	0.25-8	1	2
	Erythromycin	0.25-16	1	2
	Levofloxacin	1-8	2	4
	Doxycycline	0.03-0.5	0.25	0.5

## Discussion

- CEM-101 showed significant activity (MIC<sub>90</sub> ≤1 mg/L) against categorized *Streptococcus pneumoniae* strains, including strains that were resistant to macrolides (*erm* B or *mefE* genotype) or quinolones.
- Against erythromycin-resistant (*erm* B genotype) *S. pneumoniae*, CEM-101 was significantly superior to the antibiotics tested: telithromycin, azithromycin and erythromycin, doxycycline and levofloxacin.
- When *S. pneumoniae* ciprofloxacin-resistant (*gyrA* and *parC* genotype) strains were treated with CEM-101, this new macrolide exerted greater activity (MIC<sub>90</sub> 0.25 mg/L) and was superior to doxycycline. This observation was not seen with the other tested macrolides.
- The activity (MIC<sub>90</sub> 4 mg/L) of CEM-101 was clearly superior to all macrolides tested (MIC<sub>90</sub> ≥16 mg/L) against erythromycin-resistant *H. influenzae* (*erm*A, B, C genotype).

## Conclusion

- CEM-101 shows a broad spectrum of activity against the most commonly isolated resistant strains of *S. pneumoniae* or *H. influenzae* isolated from respiratory tract infections.
- With favorable pharmacokinetics in humans, CEM-101 should be a valuable oral compound for the treatment of upper or lower respiratory tract infections caused by *S. pneumoniae* or *H. influenzae* that are resistant to standard oral macrolides or quinolones.
- Clinical studies should undertaken to evaluate the *in vivo* effectiveness of this new antimicrobial agent.

## References

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