



## **SENSIBILITY OF CLINICAL ISOLATES TO LEVOFLOXACIN UNDER THE CONDITIONS OF INCREASING ANTIMICROBIAL RESISTANCE**

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### **ABSTRACT**

The appearance of comparatively new diseases, caused by multi-resistant microbial species and their toxins, as well as the increase in the population of patients at risk imposes the application of contemporary approach when choosing antimicrobial therapy. Suitable medicine which meets the modern needs of antimicrobial therapy is Levofloxacin. The purpose of this recent investigation is to determine the antimicrobial resistance of the most frequent clinical isolates to Levofloxacin. The resistance of 183 isolates of Gram (+) and Gram (-) bacteria ( *S. aureus* , *S. pneumoniae* , *S. agalactiae* , *Enterococcus* spp. , family *Enterobacteriaceae* , *P. aeruginosa* and *Acinetobacter* spp.) has been investigated , using DDM and BDM . One of the 25 investigated *S. aureus* strains has shown resistance to Levofloxacin ,while *S. pneumoniae* and *S. agalactiae* have not.The half of the tested *Enterococcus* spp. strains have been levofloxacin resistant. Among 119 investigated Gram (-) bacteria, 41 have shown resistance to Levofloxacin, (mainly due to *Enterobacteriaceae*, producing Extended Spectrum Beta Lactamases and multi –resistant *Acinetobacter* spp.)

**Key words:** Levofloxacin ,Gram (+) cocci, *Enterobacteriaceae*, *P. aeruginosa*, *Acinetobacter* spp.

### **INTRODUCTION**

Despite the serious success in antibiotic treatment of the infections, the antimicrobial therapy is still of great challenge for medicine. The application of contemporary approach in the choice of antimicrobial medicaments depends on various factors:

1. The appearance of new diseases, caused by new bacterial species and their toxins: Legionellosis, caused by *L. pneumophila*; Lyme Borreliosis, caused by *B. burgdorferi*; necrotising fasciitis, caused by *S. pyogenes*; TSS - Toxic Shock Syndrome, caused by toxigenic *S. aureus*, producing TSST-1 (Toxic Shock Syndrome Toxine 1); antibiotic associated colitis due to *C. difficile*-enterotoxin.

2. The wider spreading of multi-resistant microbial species such as: *S. aureus*-vancomycin – intermediate; *S.pneumoniae* - penicillin-resistant; *Enterococcus faecium*-oxazolidinones- and vancomycin-resistant, Gram (+) cocci with MLS (macrolides-

lincosamides-streptogramins)-types of resistance, members of the family *Enterobacteriaceae* producing Extended Spectrum Beta Lactamases (ESBLs), Gram (-) nonfermentative bacteria with multi-resistance.

3. The increase in the number of the patients at risk predisposes on the one hand to infections with multi-resistant microorganisms, on the other to severer course and higher mortality rate.

4. Medical practice has to dispose with appropriate peroral and parenteral drugs for empirical therapy, as well as with such for the continuing treatment.

5. For the treatment of infections, caused by multi-resistant microbial species, there are only a few numbers of antimicrobial medicaments.

- Against Gram (+) species: glycopeptides, oxazolidinones ,streptogramins etc.
- Against Gram (-) species : cephalosporins, aminoglycosides etc.

Modern antimicrobial medicine, which meets the requirements of contemporary approach in antimicrobial therapy, is Levofloxacin. It has powerful bactericidal effect, wide antibacterial spectrum, excellent penetration in pulmonary

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tissue and sinuses, high intercellular concentrations. The chemical formula of Levofloxacin is with low potential for developing resistance in bacterial species.

Levofloxacin inhibits the enzymes which catalyse the process of replication of DNA - deoxyribonucleic acid - gyrase and topoisomerase IV. The inhibition of these enzymes leads to bacterial death.

The spectrum of activity of Levofloxacin includes a great number of Gram (+) and Gram (-) microorganisms such as *Haemophilus influenzae*, *Moraxella catarrhalis*, *Enterobacter* spp., *Escherichia coli*, *Klebsiella* spp., *Proteus* spp., *Serratia* spp., *Pseudomonas aeruginosa*, *Stenotrophomonas maltophilia*, *Acinetobacter* spp.

The medicine is suitable for the treatment of widely spread intercellular (atypical) microorganisms: *Chlamydia pneumoniae*, *Mycoplasma pneumoniae*, *Ureaplasma urealyticum*, *Legionella* spp. (1-3).

Levofloxacin is the first choice drug of empirical therapy of patients at risk with pulmonary infections (smokers, immunosuppressed and other), urogenital infections, wound infections and etc. Thereby this medicine is one of the most appropriate and often used antimicrobial drug in medical practice (4-7).

## PURPOSE

The aim of this recent investigation is to determine the antimicrobial resistance of the most frequent clinical isolates to Levofloxacin during one year period.

## MATERIALS AND METHODS

There were investigated totally 183 strains; 64 Gram (+) and 119 Gram (-) bacteria, isolated from infections of upper and lower respiratory tract, urogenital and surgical infections. The antimicrobial susceptibility of the isolates was determined via DDM (Disk Diffusion Method) and BDM (Broth Dilution Method). The interpretation of the results was done according to CLSI (Clinical Laboratory Standard Institute) 2009.

## RESULTS

The results of the antimicrobial resistance of Gram (+) isolates is shown on **Table 1**.

From all of the investigated 65 Gram (+) isolates, 15 were resistant to Levofloxacin. Resistant to Levofloxacin was only 1 singular isolate of *S. aureus*, the same was also multi-resistant -MRSA and MLS. Among the isolates of *S. pneumoniae* and *S. agalactiae* there were not resistant to Levofloxacin, although some strains were multi-resistant among them: five isolates PNSSP, three isolates of *S. pneumoniae* and two of *S. agalactiae* with MLS-resistance. The most of the tested *Enterococcus* species (eight from fifteen), were resistant to Levofloxacin, which is usual for *Enterococci*. Among Levofloxacin resistant isolates of *Enterococci*, two were multi-resistant, while the others were resistant only to fluoroquinolones.

Among Gram (-) bacteria there was considerably higher resistance to Levofloxacin (**Table 2**). Among 119 isolates, being tested, 41 were Levofloxacin resistant. These resistant strains were mainly multi-drug resistant *P.aeruginosa* and *Acinetobacter* spp. and *Enterobacteriaceae*, producing ESBL<sub>s</sub>.

**Table 1.** Resistance to Levofloxacin of Gram (+) isolates of different phenotypes:

	N	Levofloxacin-R
<i>S. aureus</i> -Total	25	1
<i>S. aureus</i> -MSSA	24	0
<i>S.aureus</i> -MRSA	1	1
<i>S. aureus</i> -MLS	4	1
<i>S. pneumoniae</i> -Total	15	0
<i>S. pneumoniae</i> PNSSP	5	0
<i>S. pneumoniae</i> -MLS	3	0
<i>S. agalactiae</i> -Total	10	0
<i>S. agalactiae</i> -MLS	2	0
<i>Enterococcus</i> spp.	15	8
<i>Enterococcus</i> spp.-MDR	2	2

MSSA-methicillin- sensitive *S. aureus*

MRSA- methicillin-resistant *S. aureus*

MLS – resistance to macrolides, lincosamides and streptogramins

PNSSP – penicillin-nonsensitive *S. pneumoniae*

MDR- multi-drug resistant

## CONCLUSIONS

- Under the conditions of constantly increasing antimicrobial resistance Levofloxacin keeps its activity against the most of the clinically significant Gram (+) microorganisms such as *S. aureus*, *S. agalactiae*, *S. pneumoniae* (including

PNSSP, MLS) and family Enterobacteriaceae.

- Resistance to Levofloxacin more often display bacteria with multi-drug resistance such as *Enterococcus* spp. , Enterobacteriaceae – ESBL<sub>s</sub> (+), *P. aeruginosa*, *Acinetobacter* spp.

**Table 2.** Resistance to Levofloxacin of Gram ( - ) isolates of different phenotypes of resistance

Mechanism/phenotype of resistance	N	Levofloxacin-R
Enterobacteriaceae - total	84	22
Enterobacteriaceae ESBL <sub>s</sub> (+)	19	11
<i>P. aeruginosa</i> - total	20	10
<i>P. aeruginosa</i> - MDR	7	7
<i>Acinetobacter</i> spp.	15	12
<i>Acinetobacter</i> spp. - MDR	12	11

ESBL<sub>s</sub> – Extended spectrum β-lactamase

MDR - Multi-drug resistant

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