

Activity of finafloxacin (FNX), a fluoroquinolone with enhanced activity at acid pH, against intracellular *Legionella pneumophila*: comparison with azithromycin, clarithromycin, telithromycin, ciprofloxacin and moxifloxacin

P1139

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Abstract

Background and aim

Results

Background:
L. pneumophila invades macrophages and multiplies in moderately acidic vacuoles (Sturgill-Koszycki & Swanson, *J Exp Med* 2000, 192:1261-72). Macrolides and quinolones are commonly recommended treatment for LP infections. In this context, our aim was to examine the activity FNX (a novel fluoroquinolone exhibiting increased activity under acidic conditions [Higgins et al., *AAC* 2010, 54:1613-5]) against intracellular LP.

Methods:

L. pneumophila

ATCC 33153

and THP-1

cells

were

used

. MICs

and

C_{max}

were

measured

in

α-ketoglutarate

buffered

yeast

extract

(pH

6.9)

after

48

h

incubation

.

Non-phagocytized

bacteria

were

eliminated

by

incubation

in

Phosphate

Buffered

Saline

(PBS)

supplemented

with

50

mg/L

gentamicin

(1

h)

MIC:

0.25

mg/L

and

4

successive

washings

with

PBS

.

Cells

were

then

transferred

to

fresh

culture

medium

containing

FNX

and

comparators

covering

a

wide

range

of

concentrations

to

obtain

full

concentration-dependent

effects

.

After

48

h

cells

were

harvested

washed

with

PBS

,

lysed

and

used

for

enumeration

of

CFU

and

assay

of

cell

protein

.

Data

were

used

to

determine

the

apparent

static

concentration

(Cs)

of

each

antibiotic

and

its

activity

at

C_{max}

.

Results

and

Methods

are

described

elsewhere

.

Conclusions:

Compared

to

macrolides

,

quinolones

appear

more

effective

against

intracellular

LP

.

Amongst

them

, FNX

shows

the

greatest

activity

at

clinically-relevant

concentrations

,

perhaps

in

relation

to

its

improved

activity

at

acidic

pH

.

Electron

microscopy

.

Cells

were

infected

as

described

above

, except

that

the

initial

inoculum

was

increased

to

20

bacteria

per

macrophage

.

Sample

handling

was

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