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Potent activity of BAY 73-7388, a novel aminomethylcycline, against susceptible and resistant Gram-positive and Gramnegative organisms

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Objectives: BAY 73-7388 is the first of a new class of antibiotics, the aminomethylcyclines, which evolved from the tetracycline (TET) family. BAY 73-7388 has potent activity against antibiotic susceptible and resistant Gram-positive and Gram-negative pathogens. The present study compared the activity *in vitro* of BAY 73-7388 and 10 other agents including vancomycin (VAN), linezolid (LIN), levofloxacin (LVX) and TET against recent clinical isolates including MRSA, VAN-resistant *Enterococcus faecium* (Efa VRE), *Enterococcus faecalis* (Ef), penicillin-resistant *Streptococcus pneumoniae* (Spn PENR), Groups A and B beta-haemolytic streptococci (BHS), *Escherichia coli* (Ec), and other pathogens. Potential microbiological interactions between BAY 73-7388 and other antibiotics were also assessed.

Methods: Microdilution MIC tests were performed according to NCCLS guidelines. The activity of BAY 73-7388 in the presence of other antibacterial agents was assessed using standard chequerboard MIC methods. TET-resistance determinants were identified using multiplex PCR.

Results: Susceptibility in vitro (MIC90 mg/L) for selected agents is shown in the table below.

	MIC 90 (mg/L)						
Strains	BA Y 73- 7388	VA N	LI N	LV X			
MRSA (<i>n</i> = 39)	0.5	1.0	2.0	32.0			
Efa VRE $(n = 19)$	0.5	>64. 0	2.0	>64. 0			
Ef (<i>n</i> = 31)	0.5	2.0	2.0	32.0			
Spn PENR $(n = 23)$	0.06	0.25	1.0	1.0			
BHS (<i>n</i> = 48)	0.25	0.5	1.0	0.5			
Ec (<i>n</i> = 23)	2.0	NA	NA	4.0			
,							

Conclusions: BAY 73-7388 has potent activity *in vitro* against a range of common pathogens, including those resistant to currently available antibiotics: most notably MRSA, VRE and penicillin-resistant *S. pneumoniae*. Chequerboard studies *in vitro* demonstrate BAY 73-7388 does not affect, and is not affected by, the activity of other antibiotics.

(BAY 73-7388 was discovered by Paratek Pharmaceuticals Inc., Boston, MA, and designated PTK 0796.)

POTENT ACTIVITY OF BAY 73-7388, A NOVEL AMINOMETHYLCYCLINE, AGAINST SUSCEPTIBLE AND RESISTANT GRAM-POSITIVE AND GRAM-NEGATIVE ORGANISMS



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Table 2. Comparative in vitro activity of BAY 73-7388

1.0-8.0 ≤0.06>64.0 0.5-32.0 ≤0.06>64.0 ≤0.06 64.0 0.5>64.0

2.0>64.0

Table 3. In vitro activity of BAY 73-7388 combined

Test

Gertamicin

(mg/L)

0.5 1.5

0.03 2.0 0.5

MIC BAY 73-7388

(mg/L)

0.5

1.0

0.125

FICI interpretation: Synergy <0.5; No interaction <0.5-4.0; Antagonism >4.0.

MIC (mg/L)

0.5 ≤0.06 ≤0.06 2.0 2.0 2.0

vs Escherichia coli, Klebsiella pneumoniae, and

Haemophilus influenzae

BAY 73-7388

Ampicillin Cefetaxime Gentamicin Ciproflusacin Levoflosacin Tetracycline Minosycline Doxycycline

BAY 73-7388 Celotaxime Gentamicin Ciprofloxacin Lovofloxacin Tetracycline Minocycline Doxyccline

BAY 73-7388

Organism

Strain

E. coli 225

E. coll ATOC 25922

E. faecalis ATCC 29212

Abstract

Objectives:

BAY 73-7388 is the lead compound of a new class of antibiotics, the aminomethylcyclines, which evolved from the tetracycline (TET) family, BAY 73-7388 has potent activity against antibiotic-susceptible and resistant gram-positive and gram-negative pathogens. The present study compared the activity in vitro of BAY 73-7388 and 10 other agents including vancomycin (VAN), linezolid (LZD), levofloxacin (LVX) and TFT against recent clinical isolates including methicillin-resistant. Staphylococcus aureus (MRSA), VAN-resistant Enterococcus faecium (Efa VRE), Enterococcus faecalis (Ef), penicillin-resistant Streptococcus pneumoniae (Spn PENR), Groups A and B beta-haemolytic streptococci (BHS), Escherichia coli (Ec), and other pathogens. Potential microbiological interactions between BAY 73-7388 and other antibiotics were also assessed.

Methods:

Microdilution MIC tests were performed according to NCCLS guidelines. The activity of BAY 73-7388 in the presence of other antibacterial agents was assessed using standard chequerboard MIC methods. TET-resistance determinants were identified using multiplex polymerase chain reaction (PCR).

Susceptibility in vitro (MIC 90 mg/L) for selected agents is shown in the Table below.

Conclusions:

BAY 73-7388 has potent activity in vitro against a range of common pathogens, including those resistant to currently available antibiotics: most notably MRSA, VRE and penicillin-resistant S.pneumoniae. Chequerboard studies in vitro demonstrate BAY 73-7388 does not affect, and is not affected by, the activity of other antibiotics.

	MIC ₉₀ (mg/L)						
Strains	BAY 73-7388	VAN	LZD	LVX			
MRSA (n-39)	0.5	1.0	2.0	32.0			
Efa VRE (n=19)	0.5	>64.0	2.0	>64.0			
Ef (n=31)	0.5	2.0	2.0	32.0			
Spn PENR (n=23)	≤0.06	0.25	1.0	1.0			
BHS (n-48)	0.25	0.5	1.0	0.5			
Ec (n=23)	2.0	NA	NA	4.0			

Figure 1. Structure of BAY 73-7388.

(BAY 73-7388 was discovered by Paratek Pharmaceuticals Inc., Boston, MA, and designated PTK 0796.)

Introduction

- BAY 73-7388, 7-dimethylamino,9-(2,2-dimethylpropyl)-aminomethylcycline is the first of a new class of antibiotic, the aminomethylcyclines (Figure 1)
- In this study, the activities of BAY 73-7388 and comparator compounds were determined against a variety of aerobic gram-positive and gram-negative bacteria including strains resistant to currently available classes of antibiotics
- The activity of BAY 73-7388 was determined alone and in combination with antibiotics having differing mechanisms of action

Methods

Strains

- · Recent clinical isolates obtained from 7 laboratories within the United States
- Appropriate quality control strains were included in each test

MIC Determination

- By current NCCLS-recommended microdilution method¹
- 1. Cation-adjusted Mueller Hinton broth for staphylococci, enterococci, and gram-negative rods
- 2. Cation-adjusted Mueller Hinton broth supplemented with 5% horse blood for streptococci
- 3. Haemophilus test medium for Haemophilus

- Inoculum: Organisms were grown (or direct inoculum for fastidious organisms) to a 0.5 McFarland standard. Turbidity was measured using a Microscan turbidity meter.
- Incubation at 35°C for 18 to 24 hours depending on the organisms tested

Chequerboard MIC

By standard broth microdilution method²

Fractional inhibitory concentration index was calculated as FIC_A + FIC_B where FIC represents the minimum concentrations that inhibited growth for compounds A and B/the MIC of A or B alone.

Table 1. Comparative in vitro activity of BAY 73-7388 vs aerobic gram-positive bacteria

				MIC (mg/L						MIC (mg/	
Organism	N	Compound	Range	MIC ₅₀	MIC ₉₀	Organism	N	Compound	Range	MIC ₅₀	MIC ₉₀
S. aureus	55	BAY 73-7388	≤0.06-1.0	0.125	0.5	E. faecalls	31	BAY 73-7388	0.1255	0.25	0.5
	•	Vancomycin	0.25-2	0.5	1	E. IDOUND	01	Vancomycln	0.5-8	1	2
		Linezolid	0.5-2.0	2	2			Linezolid	1.0-4.0	1	2
		Levofloxacin	≤0.06->64.0	4	32			Levoffoxacin	0.5-64	1	32
		Azithromycin	0.25->64.0	>64.0	>64.0			Azithromycin	1.0->64.0	8	>64.0
		Clindamycin Cefotaxime	≤0.06->64.0 1.0->64.0	0.125 32	>64.0 >64.0			Clindamycin Tetracycline	2.0->64.0 0.125->64.0	32 32	>64.0 64
		Tetracycline	≤0.06-64.0	0.125	64			Minocycline	0.125-564.0	8	16
		Minocycline	≤0.06-16.0	0.125	8			Doxycycline	≤0.06-16.0	4	16
		Daxycycline	≤0.06-8.0	≤0.06	8	E. faecalis	3		0.25-0.5	NA.	NA.
S. aureus: Methicillin-resistant	39	BAY /3-/388	0.125-1.0	0.25	0.5	E. raecans Multiresistant	3	BAY 73-7388 Vancomycin	0.25-0.5	NA NA	NA NA
	35	Vancomycin	0.25-2.0	0.25	1	Multiresistant		Linezolid	1	NA.	NA.
inconciliii reaseant		Linezolid	0.5-2.0	2	2			Levofloxacin	16.0-64.0	NA.	NA.
		Levofloxacin	0.5->64.0	8	32			Azithromycin	>64.0	NA	NA
		Azithromycin	0.5->64.0	>64.0	>64.0			Clindamycin	>64.0	NA	NA
		Clindamycin	≤0.06->64.0	>64.0	>64.0			Tetracycline	32.0-64.0	NA	NA
		Cefotaxime Tetracycline	4.0->64.0 ≤0.06-64.0	>64.0 0.25	>64.0 64			Minocycline Doxycycline	8.0-16.0 4	NA NA	NA NA
		Minocycline	≤0.06-16.0	0.25	8						
		Daxycycline	≤0.06-8.0	0.125	8	S. pneumon/ae	41	BAY 73-7388	≤0.06-0.25	≤0.06	0.125
								Vancomycin	≤0.06-0.5	0.25	0.25
S. aureus:	16	BAY 73-7388	≤0.06-0.25	0.125	0.125			Linezolid	0.25-2.0	1.0	1.0
Methicillin-sensitive		Vancomycin Linezolid	0.25-0.5 1.0-2.0	0.5 1	0.5 2			Levofloxacin Azithromycin	0.25-1.0 ≤0.06->64.0	0.5 2.0	1.0 >64.0
		Levofloxacin	<0.06-4.0	0.125	0.125			Clindamycin	≤.06->64.0	<0.06	>64.0
		Azithromycin	0.25-32.0	0.5	0.5			Cefotaxime	≤0.06-8.0	1.0	2.0
		Clindamycin	≤0.06-0.125	≤0.06	0.125			Penicillin	0.8-8.0	2.0	4.0
		Cefotaxime	1.0-2.0	2	2			Tetracycline	≤0.06-64.0	16.0	32.0
		Tetracycline	<0.06-16.0	< 0.06	0.125			Minocycline	<0.06-8.0	2.0	8.0
		Minocycline	≤0.06-0.125 ≤0.06-1.0	≤0.06 ≤0.06	0.125 ≤0.06			Doxycycline	≤0.06-4.0	2.0	4.0
		Darycycline				S. pneumon/ae	23	BAY 73-7388	≤0.06	≤0.06	≤0.06
S. aureus:	10	BAY 73-7388	0.25-0.5	0.5	0.5	Penicillin-resistant		Vancomycin	0.125-0.25	0.25	0.25
Multiresistant		Vancomycin	0.5-1.0	1	1			Linezolid	0.5-2.0	1.0	1.0
		Linezolid Levofloracin	0.5-2.0 8.0-32.0	1 8	2 32			Levofloxacin Azithromycin	0.5-1.0 ≤0.06->64.0	0.5 4.0	1.0 >64.0
		Azithromycin	>64.0	>64.0	>64.0			Clindamycin	<0.06->64.0	<0.06	>64.0
		Clindamycin	>64.0	>64.0	>64.0			Cefotaxime	0.5-8.0	10	8.0
		Cefotaxime	32.0-64.0	>64.0	>64.0			Penicillin	2.0-8.0	4.0	8.0
		Tetracycline	32.0->64.0	>64.0	>64.0			Tetracycline	≤0.06-64.0	32.0	32.0
		Minocycline	2.0-16.0	8	8			Minocycline	0.125-8.0	8.0	8.0
		Daxyoyaline	2.0 8.0	8	8			Doxyoyeline	≤0.06 4.0	4.0	4.0
E. faecium	24	BAY 73-7388	0.125-0.5	0.25	0.5	S. pneumon/ae	18	BAY 73-7388	≤0.06	≤0.06	≤0.06
		Vancomycin	0.5->64.0	>64.0	>64.0	Multiresistant		Vancomycin	0.125-0.25	0.125	0.25
		Linezolid	0.5-4.0	2 64	2			Linezolid	0.5-1.0	1.0	1.0
		Levofloxacin Azithromycin	1.0->64.0 4.0->64.0	>64.0	>64.0 >64.0			Levofloxacin Azithromycin	0.5-1.0 2.0->64.0	0.5 >64.0	1.0 >64.0
		Clindamycin	≤0.06>64.0	>64.0	>64.0			Clindamycin	≤0.06.>64.0	>64.0	>64.0
		Tetracycline	0.125 >64.0	32	64			Cefotaxime	0.5 8.0	1.0	8.0
		Minocycline	0.125-32.0	8	16			Penicillin	2.0-8.0	4.0	8.0
		Daxycycline	≤0.06-16.0	2	8			Tetracycline	16.0-64.0	32.0	32.0
E faecium	19	BAY 73-7388	0.125-0.5	0.25	0.5			Minocycline	4.0-8.0	8.0	8.0
Vancomycin-resistant		Vancomycin	64.0->64.0	>64.0	>64.0			Doxycycline	2.0-4.0	4.0	4.0
		Linezolid	0.5-4.0	2	2	S. pyogenes	30	BAY 73-7388	≤0.06-0.5	0.125	0.25
		Levofloxacin	1.0->64.0	64	>64.0			Vancomycin	0.25	0.25	0.25
		Azithromyoin	>64.0	>64.0	>64.0			Linozolid	0.5 1.0	1.0	1.0
		Clindemycin	>64.0 0.125->64.0	>64.0 32	>64.0 64			Levofloxacin Azithmorwcin	0.25-1.0 ≤0.06→64.0	0.25 ≤0.06	1.0
		Tetracycline Minocycline	0.125-964.0	8	16			Clindamycin	≤0.06->64.0	≤0.06	≤0.06
		Dusycycline	≤0.06-8.0	2	4			Cefotaxime	≤0.06	≤0.06	≤0.06
								Tetracycline	≤0.06-64.0	≤0.08	64.0
E. faecium Multimeistant	12	BAY 73-7388 Vancomycin	0.125-0.5 >64.0	0.25 >64.0	0.5 >64.0			Minocycline	0.125-8.0	0.25	8.0
mundresistant		Vancomycin Linezolid	0.5 2.0	>61.0 1	N64.0 2			Doxycycline	≤0.06-8.0	≤0.06	8.0
		Levofloxacin	8.0->64.0	32	>64.0	S. agalactics	18	BAY 73-7388	≤0.06-0.25	0.125	0.125
		Azithromycin	>64.0	>64.0	>64.0			Vancomycin	0.125-0.5	0.25	0.5
		Clindamycin	>64.0	>64.0	>64.0			Linezolid	1.0-1.0	1.0	1.0
		Tetracycline	32.0->64.0	32	>64.0			Levofloxacin	0.125-0.5	0.5	0.5
		Minocycline	4.0-16.0 2.0-8.0	8 2	16 4			Azithromycin Clindamycin	≤0.06-8.0 ≤0.06	≤0.06 ≤0.06	0.125 ≤0.06
		Darycycline	2.0-8.0	2	4			Clindamyoln Cefotaxime	≤0.06 ≤0.06	≤0.06 ≤0.06	≤0.06 ≤0.06
								letracycline	≤0.06-64.0	32.0	64.0
								Minocycline	0.125-32.0	16.0	16.0
								Doxycycline	≤0.06-16.0	8.0	8.0

Results

Gram-Positive Aerobic Bacteria (Table 1)

- Methicillin-sensitive Staphylococcus aureus (MIC₉₀ = 0.125 mg/L)
- Methicillin-resistant Staphylococcus aureus (MIC₉₀ = 0.5 mg/L)
- Methicillin-resistant and multiresistant (multiR) Staphylococcus aureus (MICoo = 0.5 mg/L)
- Enterococcus faecium, including vancomycin-resistant and vancomycin-resistant, multiR strains
- Enterococcus faecalis, including vancomycin-resistant and vancomycin-resistant, multiR strains
- $(MIC_{QO} = 0.5 \text{ mg/L})$ Streptococcus pneumoniae including penicillin-resistant and penicillin-resistant, multiR strains
- $(MIC_{oo} \leq 0.06 \text{ mg/L})$
- Streptococcus pyogenes (MIC₉₀ = 0.25 mg/L)
- Streptococcus agalactiae (MIC₉₀ = 0.125 mg/L)
- BAY 73-7388 was the most active compound tested against all gram-positive bacteria

Select Gram-Negative Aerobic Bacteria (Table 2)

- . BAY 73-7388 inhibited all strains of

Chequerboard (Table 3)

· No interactions were observed between BAY 73-7388 and any of the antibiotics tested

- Methicillin-resistant Staphylococcus aureus
- · Penicillin-resistant Streptococcus pneumoniae
- · Escherichia coli
- Klebsiella pneumoniae

BAY 73-7388 is active in vitro against organisms which are resistant to multiple classes of antibiotics

- Methicillin-resistant Staphylococcus aureus and vancomycin-resistant Enterococcus also resistant to tetracyclines, levofloxacin.
- Penicillin-resistant Streptococcus pneumoniae also resistant to tetracyclines, cefotaxime.

BAY 73-7388 shows no interaction with ampicillin. cefotaxime.ciprofloxacin.clindamycin.gentamicin. rifampin, or vancomycin

- . BAY 73-7388 inhibited all strains of
- $(MIC_{oo} = 0.5 \text{ mg/L})$

- Escherichia coli (MIC_{po} = 2.0 mg/L)
- Klebsiella pneumoniae (MICon = 4.0 mg/L)
- Haemophilus influenzae (MICon = 2.0 mg/L)

Conclusions

BAY 73-7388 is highly active in vitro against a broad spectrum of pathogens, most notably:

- · Vancomycin-resistant Enterococcus

- Haemophilus influenzae

with other antimicrobial agents including:

- azithromycin, and clindamycin
- azithromycin, and clindamycin

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