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Activity of Fosfomycin against Gram-Negative Baseline Bacterial Isolates from Patients in a Phase 3 Complicated Urinary Tract Infection Trial (ZEUS) D SHORTRIDGE¹, RE MENDES¹, LN WOOSLEY¹, D SKARINSKY², EJ ELLIS-GROSSE², RK FLAMM¹ ¹JMI Laboratories, North Liberty, Iowa, USA; ²Zavante Therapeutics, Inc., San Diego, California, USA

Abstract

Background: ZTI-01 (fosfomycin, FOS, for injection) is an investigational antibiotic under US development to treat complicated urinary tract infections (cUTI). FOS is different than other antimicrobials in that it is thought to inhibit an early step in cell wall synthesis via covalent binding to MurA. FOS demonstrated broad in vitro activity against gram-negative (GN) and -positive (GP) bacteria, including organisms containing extended-spectrum β -lactamases (ESBL). In this study, we investigated the activity of FOS against GN clinical isolates from the phase 3 cUTI trial (ZEUS).

Methods: FOS was tested against 465 GN baseline clinical isolates collected in the cUTI clinical trial. Isolates were susceptibility (S) tested against comparator agents by reference broth microdilution and FOS by reference agar dilution (25 µg/mL glucose-6-phosphate supplementation). Existing FDA FOS breakpoints of $\leq 64 \text{ mg/L}$ for the oral formulation were used for comparative assessments. Screen-positive ESBL Enterobacteriaceae isolates were characterized by whole genome sequencing and analysis for the presence of known β -lactamase genes.

Results: FOS was very active against *Enterobacteriaceae* (MIC_{50/90}, 1/16 mg/L; 96.4% \leq 64 mg/L). For 329 *E. coli*, 100.0% were S to FOS (MIC_{50/90}, 1/1 mg/L), and for 62 Klebsiella pneumoniae the FOS MIC_{50/90} values were 16/128 mg/L (87.1% $\leq 64 \text{ mg/L}$). The FOS MIC_{50/90} values for 14 *Enterobacter cloacae* complex and 20 Proteus mirabilis were 16/128 and 2/32 mg/L, respectively. For 21 Pseudomonas aeruginosa, higher FOS MIC_{50/90} were observed, 64/256 mg/L. Regarding the characterized ESBL-containing isolates, the FOS MIC_{50/90} for 49 *E. coli* were 1/2 mg/L, for 28 K. pneumoniae were 16/128 mg/L, and for 3 P. mirabilis the MIC₅₀ was 2 mg/L. The most common ESBL was CTX-M (42/49 E. coli and 27/28 K. pneumoniae), with CTX-M-15 (belonging to Group 1) the most common variant. Carbapenemase genes, NDM-1 and OXA-48, were detected in 2 K. pneumoniae isolates, which had FOS MICs of 8 and 64 mg/L, respectively.

Conclusions: FOS demonstrated broad-spectrum activity against baseline clinical isolates in a phase 3 cUTI trial, including those with ESBL or carbapenemases. FOS merits further study in infections where resistant GN may occur. Potentially introducing an IV form would warrant a reassessment of susceptibility breakpoints given the current oral formulation's bioavailability limitations.

Introduction

- ZTI-01 (fosfomycin, FOS, for injection) is under US development to treat complicated urinary tract infections (cUTI)
- Fosfomycin has been used for over 40 years to treat a variety of serious infections
- FOS is different than other antimicrobials in that it is thought to inhibit an early step in cell wall synthesis via covalent binding to MurA
- FOS has demonstrated broad in vitro activity against gram-negative (GN) and -positive (GP) bacteria, including organisms containing extended-spectrum β-lactamases (ESBL)
- In this study, we investigated the activity of FOS against GN clinical isolates from the phase 3 cUTI trial (ZEUS)

Materials and Methods

• FOS was tested against 465 GN baseline clinical isolates collected in the cUTI clinical trial

Table 1 Antimicrobial activity of fosfomycin agar tested against the main organisms and organism groups of isolates (mg/L)

Organism/organism g Enterobacteriaceae (Carbapenem-resistant Escherichia coli (329) Non-ESBL-phenotype ESBL-phenotype (54) Klebsiella pneumoniae (´ Non-ESBL-phenotype ESBL-phenotype (33) Proteus mirabilis (20) Non-ESBL-phenotype ESBL-phenotype (3) Enterobacter cloacae s Pseudomonas aerugin

Acinetobacter baumann complex (2)

- comparative assessments

- FOS was very active against *Enterobacteriaceae* (MIC_{50/90}, 1/16 mg/L; 96.4% ≤64 mg/L)
- (Table 1

a activity of tostomychi agai testeu against the main organisms and organism groups of isolates (mg/L)																
group (no. of isolates)		No. of isolates at MIC (mg/L; cumulative %)											MIC ₅₀	MIC ₉₀		
	≤0.12	0.25	0.5	1	2	4	8	16	32	64	128	256	512	>	50	90
440)	0 0.0	4 0.9	154 35.9	153 70.7	33 78.2	15 81.6	20 86.1	26 92.0	13 95.0	6 96.4	8 98.2	2 98.6	1 98.9	5 100.0	1	16
ant (4)						0 0.0	2 50.0	0 50.0	0 50.0	1 75.0	1 100.0				8	N/A
)	0 0.0	4 1.2	151 47.1	148 92.1	23 99.1	2 99.7	0 99.7	0 99.7	1 100.0						1	1
pe (275)	0 0.0	3 1.1	129 48.0	123 92.7	19 99.6	1 100.0									1	1
4)	0 0.0	1 1.9	22 42.6	25 88.9	4 96.3	1 98.1	0 98.1	0 98.1	1 100.0						1	2
ae (62)			0 0.0	1 1.6	0 1.6	8 14.5	14 37.1	17 64.5	11 82.3	3 87.1	5 95.2	0 95.2	1 96.8	2 100.0	16	128
pe (29)					0 0.0	6 20.7	9 51.7	6 72.4	5 89.7	1 93.1	2 100.0				8	64
3)			0 0.0	1 3.0	0 3.0	2 9.1	5 24.2	11 57.6	6 75.8	2 81.8	3 90.9	0 90.9	1 93.9	2 100.0	16	128
		0 0.0	2 10.0	2 20.0	7 55.0	2 65.0	2 75.0	2 85.0	1 90.0	1 95.0	0 95.0	0 95.0	0 95.0	1 100.0	2	32
pe (17)		0 0.0	1 5.9	2 17.6	6 52.9	2 64.7	2 76.5	2 88.2	1 94.1	1 100.0					2	32
)		0 0.0	1 33.3	0 33.3	1 66.7	0 66.7	0 66.7	0 66.7	0 66.7	0 66.7	0 66.7	0 66.7	0 66.7	1 100.0	2	N/A
e species complex (14)			0 0.0	1 7.1	1 14.3	2 28.6	2 42.9	5 78.6	0 78.6	1 85.7	1 92.9	0 92.9	0 92.9	1 100.0	16	128
inosa (21)				0 0.0	1 4.8	1 9.5	0 9.5	0 9.5	4 28.6	9 71.4	3 85.7	1 90.5	1 95.2	1 100.0	64	256
annii-calcoaceticus species										0 0.0	2 100.0				128	N/A

Shaded column represents CLSI breakpoints for E. coli for oral formulation, for information

 Isolates were susceptibility (S) tested against comparator agents by Clinical and Laboratory Standards Institute (CLSI) reference broth microdilution and FOS by reference agar dilution (with 25 µg/mL glucose-6-phosphate supplementation)

Existing FDA FOS breakpoints of ≤ 64 mg/L for the oral formulation were used for

- CLSI and EUCAST interpretive criteria were used for comparators

 Screen-positive ESBL Enterobacteriaceae isolates were characterized by whole genome sequencing and analysis for the presence of known β -lactamase genes

Results

• The most common organism isolated was *Escherichia coli* (n=329) followed by Klebsiella pneumoniae (n=62) and Pseudomonas aeruginosa (n=21)

- For 329 *E. coli*, 100.0% were S to FOS (MIC_{50/90}, 1/1 mg/L), and for 62 K. pneumoniae, the FOS MIC_{50/90} values were 16/128 mg/L (87.1% \leq 64 mg/L)

- The FOS MIC_{50/90} values for 14 *Enterobacter cloacae* complex and 20 *Proteus* mirabilis were 16/128 and 2/32 mg/L, respectively

- For 21 *P. aeruginosa*, higher FOS MIC_{50/90} values were observed (64/256 mg/L)

Table 2 Activity of fosfomycin agar and comparator antimicrobial agents when tested against 440 isolates of Enterobacteriaceae

Antimiarabial agent	MIC ₅₀	MIC ₉₀	Range		CLSI ^a	EUCAST ^a		
Antimicrobial agent	(mg/L)	(mg/L)	(mg/L)	%S	%	%R	%S	%
Fosfomycin agar	1	16	0.25 to >512				95.0	
Meropenem	≤0.06	≤0.06	≤0.06 to >8	98.9	0.2	0.9	99.1	0.5
Imipenem	0.12	0.25	≤0.06 to >8	95.9	2.3	1.8	98.2	1.4
Amikacin	2	4	≤0.5 to >64	96.8	0.5	2.7	95.9	0.9
Gentamicin	0.5	>8	≤0.06 to >8	87.5	0.7	11.8	86.1	1.4
Levofloxacin	≤0.06	>8	≤0.06 to >8	72.7	2.3	25.0	67.7	3.0
Tetracycline	2	>32	≤0.25 to >32	60.7	1.4	38.0		
Trimethoprim-sulfamethoxazole	≤0.25	>32	≤0.25 to >32	69.3		30.7	69.3	1.1
Ceftazidime	≤0.5	64	≤0.5 to >64	80.9	2.0	17.0	78.2	2.7
Aztreonam	≤0.25	>32	≤0.25 to >32	79.1	3.0	18.0	77.5	1.6
Piperacillin-tazobactam	2	16	≤0.5 to >64	90.0	3.6	6.4	88.2	1.8
Ceftriaxone	0.06	>2	≤0.015 to >2	76.8	1.6	21.6	76.8	1.6
Nitrofurantoin	16	128	4 to >128	75.2	6.4	18.4		
Nitrofurantoin a Criteria as published by CLSI 2018. CLSI fosfomycin ora			4 to >128	75.2	6.4	18.4		

Table 3 Summary of MIC results obtained against main baseline pathogens that met the MIC-based screening criteria

	Genotype		MIC (mg/L)					
Pathogen	(n)	Agent	50%	90%	Min.	Max. 32		
E. coli	All (49)	Fosfomycin	1	2	0.25			
		Piperacillin-tazobactam	2	32	1	>64		
		Aztreonam	32	>32	1	>32		
		Ceftazidime	16	>64	≤0.5	>64		
		Meropenem	≤0.06	≤0.06	≤0.06	0.12		
	CTX-M (42)	Fosfomycin	1	2	0.25	32		
		Piperacillin-tazobactam	2	32	1	>64		
		Aztreonam	32	>32	1	>32		
		Ceftazidime	16	>64	≤0.5	>64		
		Meropenem	≤0.06	≤0.06	≤0.06	0.12		
K. pneumoniae	All (28)	Fosfomycin	16	128	1	>512		
		Piperacillin-tazobactam	32	>64	1	>64		
		Aztreonam	>32	>32	8	>32		
		Ceftazidime	>64	>64	1	>64		
		Meropenem	≤0.06	8	≤0.06	>8		
CTX-M (2		Fosfomycin	16	128	1	>512		
		Piperacillin-tazobactam	16	>64	1	>64		
		Aztreonam	>32	>32	8	>32		
		Ceftazidime	>64	>64	1	>64		
		Meropenem	≤0.06	2	≤0.06	>8		
P. aeruginosa	All (11)	Fosfomycin	64	128	32	128		
		Piperacillin-tazobactam	64	64	16	>64		
		Ceftazidime	>64	>64	2	>64		
		Meropenem	4	>8	1	>8		

reundii species complex (1), Enterobacter cloacae species complex (14), Escherichia coli (329), Klebsiella oxytoca (4), K. pneumoniae (62), Morganella morganii (2), Proteus mirabilis (20), Providencia rettgeri (2), Raoultella ornithinolytica (2), Serratia marcescens (3)

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- Table 2 shows the activity of FOS and comparators against the enterics
- The least active agents were tetracycline with 60.7%S and trimethoprimsulfamethoxazole with 69.3%S (CLSI)
- The most active agents tested were FOS, imipenem, meropenem, and amikacin with ≥95%S (EUCAST)
- The most common ESBL was CTX-M (42/49 E. coli and 27/28 K. pneumoniae), with CTX-M-15 (belonging to Group 1) the most common variant
- Table 3 shows the activity of FOS and comparators for the isolates with characterized ESBL for the 3 most common organisms: *E. coli, K. pneumoniae*, and P. aeruginosa
- The FOS MIC_{50/90} values for 49 *E. coli* were 1/2 mg/L, for 28 *K. pneumoniae* were 16/128 mg/L, and for 3 P. mirabilis the MIC₅₀ value was 2 mg/L
- For 11 *P. aeruginosa* the MIC_{50/90} was 64/128 mg/L
- Carbapenemase genes NDM-1 and OXA-48 were detected in 2 K. pneumoniae isolates, which had FOS MIC values of 8 and 64 mg/L, respectively
- 1 K. oxytoca had KPC-2, with an MIC value of 8 mg/L

Conclusions

- FOS demonstrated broad-spectrum activity against baseline clinical isolates in a phase 3 cUTI trial, including those with ESBL or carbapenemases
- FOS merits further study in infections where resistant GN may occur
- Potentially introducing an IV form of FOS would warrant a reassessment of susceptibility breakpoints given the current oral formulation's bioavailability limitations

Acknowledgements

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5.0
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12.5
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19.1
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10.0
21.6

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