

# Activity of the novel extended-spectrum $\beta$ -lactamase inhibitor AAI101 in combination with cefepime towards a challenge panel of *Acinetobacter baumannii*

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## REVISED ABSTRACT

**Background:** AAI101 is a novel extended-spectrum  $\beta$ -lactamase inhibitor (BLI), active against ESBLs and a broad array of other  $\beta$ -lactamases. AAI101 in combination with cefepime (FEP) is in Phase 2 development. Infections caused by *Acinetobacter baumannii*, a pathogen endemic to the southern United States and other global regions, are very challenging to treat, and often require combination therapy. This study examined the activity of FEP/AAI101 against a challenge panel of *A. baumannii* clinical isolates enriched with OXA carbapenemase producers.

**Materials/Methods:**  $\beta$ -Lactamases in *A. baumannii* were identified by genotyping. Broth microdilution MICs and susceptibilities were obtained following CLSI methods and breakpoints (BPs), except for ceftazidime/avibactam (CAZ/AVI) where FDA *P. aeruginosa* BPs were applied. CLSI FEP BPs were used for FEP/AAI101.

**Results:** MIC<sub>90</sub> data and percentage susceptibilities (%S) for FEP/AAI101 and comparators for all *A. baumannii* isolates examined and for different *A. baumannii* subsets are shown in the table below. FEP/AAI101 was highly active against meropenem-susceptible (MPM<sup>S</sup>) isolates. FEP/AAI101 (AAI101 fixed at 8  $\mu$ g/ml) covered 67% of OXA-51 producers and 53% of OXA-58 producers, though lower susceptibilities were obtained for strains expressing OXA-23 or OXA-24/40. All OXA-51 producers contained the ISAbat1 insertion sequence upstream of bla<sub>OXA-51</sub>. Colistin (COL) was the only agent with consistently high activity against all *A. baumannii* isolates.

Group	FEP	FEP/AAI101 [4*]	FEP/AAI101 [8*]	CAZ/AVI [4*]	AMP/SUL [2+1*]	PIP/TAZ [4*]	COL
MPM <sup>S</sup> (n = 17)	MIC <sub>90</sub>	64	8	0.06	64	32	256
	%S	70.6	94.1	100	58.8	82.4	70.6
OXA-23 (n = 30)	MIC <sub>90</sub>	>128	>128	>128	128	>256	0.5
	%S	0	0	3.3	0	0	96.7
OXA-24/40 (n = 30)	MIC <sub>90</sub>	>128	>128	>128	64	>256	4
	%S	3.3	3.3	6.7	3.3	0	86.7
OXA-51 (n = 30)	MIC <sub>90</sub>	>128	>128	>128	>128	>256	0.5
	%S	0	36.7	66.7	3.3	16.7	100
OXA-58 (n = 30)	MIC <sub>90</sub>	>128	128	64	>128	64	>256
	%S	13.3	33.3	53.3	16.7	6.7	100
All (n = 137)	MIC <sub>90</sub>	>128	>128	>128	128	>256	1
	%S	12.4	27.7	40.1	13.9	16.1	8.8

AMP, ampicillin; SUL, sulbactam; PIP, piperacillin; TAZ, tazobactam  
 \*BLI at fixed concentration in  $\mu$ g/ml or ratio as indicated

**Conclusion:** Infections by this difficult pathogen often require combination therapy, of which FEP/AAI101 may be a component.

## INTRODUCTION

AAI101 is a new extended-spectrum  $\beta$ -lactamase inhibitor belonging to the penicillanic acid sulfone class (Figure 1).

In this study, the activity of cefepime/AAI101 against *Acinetobacter baumannii*, an uncommon but difficult-to-treat pathogen, was assessed using a challenge panel comprised of clinical isolates expressing predominantly one of the principal OXA  $\beta$ -lactamases responsible for carbapenem resistance in this species (OXA-23, OXA-24/40, OXA-51, OXA-58), plus some carbapenem-susceptible isolates.

## MATERIALS & METHODS

A collection of clinical isolates of *A. baumannii*, comprised of 120 strains characterised previously for the presence of OXA carbapenemases [1] plus 17 meropenem-susceptible (MPM<sup>S</sup>) isolates, were tested (Table 1). The MPM<sup>S</sup> reference strain *A. baumannii* ATCC was also included in the test panel.

o All OXA-51-producing isolates had an ISAbat1 insertion sequence upstream of the intrinsic chromosomal bla<sub>OXA-51</sub> [2].

MICs for cefepime/AAI101 [at fixed AAI101 concentrations of 4 or 8  $\mu$ g/ml] and comparator antibiotics were determined by broth microdilution using CLSI methodology [3]. AAI101 and avibactam were provided by Allegra Therapeutics SAS, and other antibacterials were purchased from commercial sources.

Susceptibility was determined according to CLSI breakpoints [4] except for:

- ceftazidime/avibactam, for which FDA breakpoints for *Pseudomonas aeruginosa* were used [5];

- cefoperazone, for which FDA breakpoints were used [6]; and

- tigecycline, for which FDA breakpoints (for *Enterobacteriaceae*) were used [7].

For comparative purposes, susceptibility of AAI101 combinations with cefepime was determined using CLSI breakpoints for cefepime alone [4], whereas cefoperazone/sulbactam susceptibility was determined using FDA breakpoints for cefoperazone alone [6].

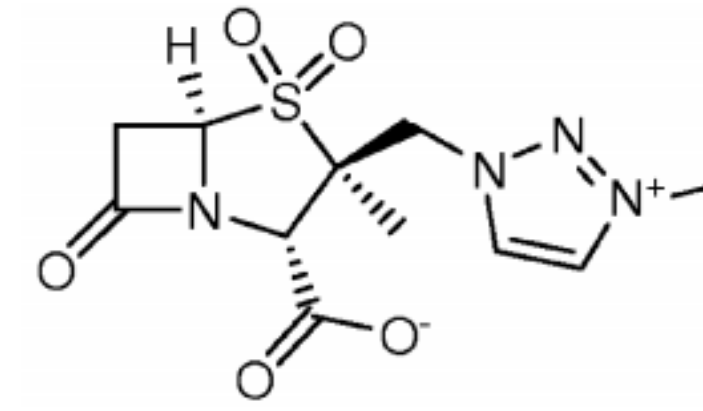


Figure 1. Chemical structure of AAI101

Table 1. Source of clinical isolates investigated

Country of origin	Meropenem-resistant isolates:				Meropenem-susceptible isolates	Total
	OXA-51	OXA-23	OXA-24/40	OXA-58		
Africa		3		3	1	7
Asia	4	3		3	2	12
Europe		17	23	13	9	62
Latin America	8	3		11	3	25
Middle East		2				2
North America	18	1	7		2	28
South Pacific		1				1
Total	30	30	30	30	17	136

Table 2. Activity of AAI101, cefepime, and AAI101 combined with cefepime against the *A. baumannii* panel

Antimicrobial (breakpoints used: S I R)	Genotype	Percentage			MIC ( $\mu$ g/ml)*	
		S	I	R	MIC <sub>50</sub>	MIC <sub>90</sub>
AAI101 (- - - -)	All (n = 137)	-	-	-	64	>128
	MPM <sup>S</sup> (n = 17)	-	-	-	4	8
	OXA-23 (n = 30)	-	-	-	64	>128
	OXA-24/40 (n = 30)	-	-	-	128	>128
	OXA-51 (n = 30)	-	-	-	16	64
OXA-58 (n = 30)	-	-	-	32	64	
Cefepime ( $\leq 8$   16   $\geq 32$ )	All (n = 137)	12.4	9.5	78.1	64	>128
	MPM <sup>S</sup> (n = 17)	70.6	17.6	11.8	4	64
	OXA-23 (n = 30)	0	0	100	>128	>128
	OXA-24/40 (n = 30)	3.3	13.3	83.3	64	>128
	OXA-51 (n = 30)	0	6.7	93.3	32	>128
OXA-58 (n = 30)	13.3	13.3	73.3	32	>128	
Cefepime/AAI101 [4 $\mu$ g/ml] ( $\leq 8$   16   $\geq 32$ )	All (n = 137)	27.7	15.3	56.9	32	>128
	MPM <sup>S</sup> (n = 17)	94.1	5.9	0	0.125	8
	OXA-23 (n = 30)	0	3.3	96.7	128	>128
	OXA-24/40 (n = 30)	3.3	20	76.7	64	>128
	OXA-51 (n = 30)	36.7	26.7	36.7	16	>128
OXA-58 (n = 30)	33.3	16.7	50	16	128	
Cefepime/AAI101 [8 $\mu$ g/ml] ( $\leq 8$   16   $\geq 32$ )	All (n = 137)	40.1	11.7	48.2	16	>128
	MPM <sup>S</sup> (n = 17)	100	0	0	$\leq 0.06$	0.06
	OXA-23 (n = 30)	0	10	90	128	>128
	OXA-24/40 (n = 30)	6.7	20	73.3	64	>128
	OXA-51 (n = 30)	66.7	6.7	26.7	4	>128
OXA-58 (n = 30)	53.3	16.7	30	8	64	

\*For cefepime/AAI101 combinations, MIC values refer to the cefepime component.

## RESULTS

Figure 2. Cumulative MIC distribution for  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations towards *A. baumannii* (n = 137)

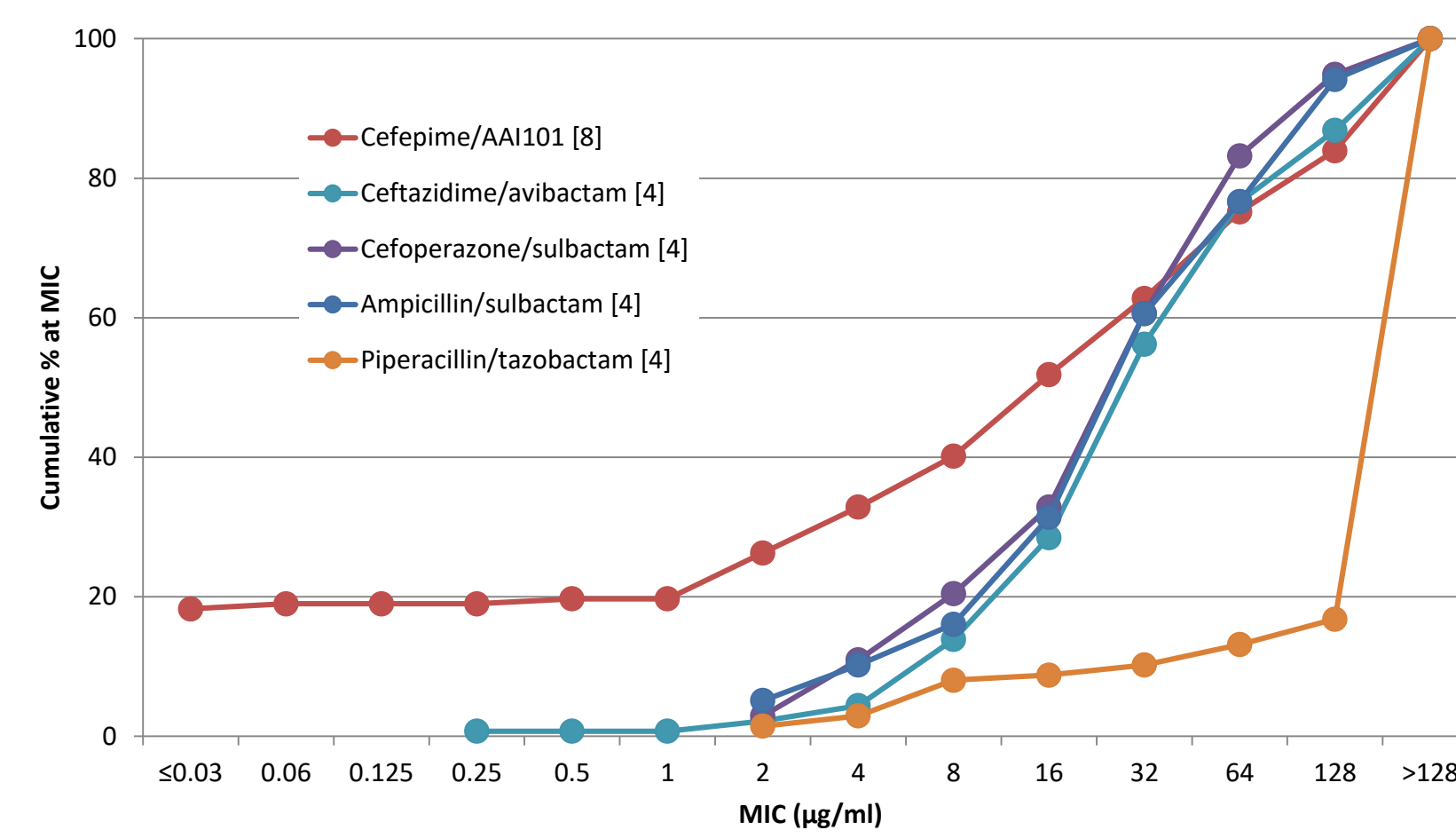


Table 3. Activity of other  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations against *A. baumannii*

Antimicrobial (breakpoints used: S I R)	Genotype	Percentage			MIC ( $\mu$ g/ml)*	
		S	I	R	MIC <sub>50</sub>	MIC <sub>90</sub>
Ampicillin/sulbactam [4 $\mu$ g/ml] ( $\leq 8$   16   $\geq 32$ )	All (n = 137)	16.1	15.3	68.6	32	128
	MPM <sup>S</sup> (n = 17)	82.4	5.9	11.8	4	32
	OXA-23 (n = 30)	0	10	90	64	128
	OXA-24/40 (n = 30)	3.3	10	86.7	32	128
	OXA-51 (n = 30)	16.7	23.3	60	32	>128
OXA-58 (n = 30)	6.7	23.3	70	32	64	
Cefoperazone ( $\leq 8$   -   $\geq 16$ )	All (n = 137)	0	0	100	>128	>128
	MPM <sup>S</sup> (n = 17)	0	0	100	64	>128
	OXA-23 (n = 30)	0	0	100	>128	>128
	OXA-24/40 (n = 30)	0	0	100	>128	>128
	OXA-51 (n = 30)	0	0	100	>128	>128
OXA-58 (n = 30)	0	0	100	>128	>128	
Cefoperazone/sulbactam [4 $\mu$ g/ml] ( $\leq 8$   -   $\geq 16$ )	All (n = 137)	20.4	-	79.6	32	128
	MPM <sup>S</sup> (n = 17)	88.2	-	11.8	4	32
	OXA-23 (n = 30)	0	-	100	64	128
	OXA-24 (n = 30)	6.7	-	93.3	32	128
	OXA-51 (n = 30)	23.3	-	76.7	32	128
OXA-58 (n = 30)	13.3	-	86.7	32	64	
Ceftazidime ( $\leq 8$   16   $\geq 32$ )	All (n = 137)	16.1	5.1	78.8	128	>128
	MPM <sup>S</sup> (n = 17)	82.4	5.9	11.8	8	128
	OXA-23 (n = 30)	10	0	90	>128	>128
	OXA-24 (n = 30)	6.7	10	83.3	64	>128
	OXA-51 (n = 30)	0	3.3	96.7	128	>128
OXA-58 (n = 30)	10	6.7	83.3	>128	>128	
Ceftazidime/avibactam [4 $\mu$ g/ml] ( $\leq 8$   -   $\geq 16$ )	All (n = 137)	13.9	-	86.1	32	>128
	MPM <sup>S</sup> (n = 17)	58.8	-	41.2	8	64
	OXA-23 (n = 30)	3.3	-	96.7	64	>128
	OXA-24 (n = 30)	6.7	-	93.3	32	64
	OXA-51 (n = 30)	3.3	-	96.7	32	>128
OXA-58 (n = 30)	16.7	-	83.3	32	>128	
Meropenem ( $\leq 2$   4   $\geq 8$ )	All (n = 137)	13.1	2.9	83.9	32	>128
	MPM <sup>S</sup> (n = 17)	100	0	0	0.5	2
	OXA-23 (n = 30)	0	0	100	64	128
	OXA-24 (n = 30)	0	0	100	128	>128
	OXA-51 (n = 30)	3.3	0	96.7	32	64
OXA-58 (n = 30)	0	13.3	86.7	16	32	
Piperacillin/tazobactam [4 $\mu$ g/ml] ( $\leq 16$   32-64   $\geq 128$ )	All (n = 137)	8.8	4.4	86.9	>256	>256
	MPM <sup>S</sup> (n = 17)	70.6	11.8	17.6	8	256
	OXA-23 (n = 30)	0	0	100	>256	>256
	OXA-24 (n = 30)	0	3.3	96.7	>256	>256
	OXA-51 (n = 30)	0	10	90	256	>256
OXA-58 (n = 30)	0	0	100	>256	>256	

\*For  $\beta$ -lactam/ $\beta$ -lactamase inhibitor combinations, MIC values refer to the  $\beta$ -lactam component.

Table 4. Activity of non- $\beta$ -lactams against *A. baumannii*

Antimicrobial (breakpoints used: S I R)	Genotype	Percentage			MIC ( $\mu$ g/ml)	
		S	I	R	MIC <sub>50</sub>	MIC <sub>90</sub>
Ciprofloxacin ( $\leq 1$   2   $\geq 4$ )	All (n=137)	9.5	2.9	87.6	>16	>16
	MPM <sup>S</sup> (n=17)	70.6	5.9	23.5	0.25	>16
	OXA-23 (n=30)	0	0	100	>16	>16
	OXA-51 (n=30)	0	0	100	>16	>16
	OXA-58 (n=30)	3.3	10	86.7	>16	>16
Colistin ( $\leq 2$   -   $\geq 4$ )	All (n=137)	96.4	-	3.6	0.5	1
	MPM <sup>S</sup> (n=17)	100	-	0	0.5	1
	OXA-23 (n=30)	96.7	-	3.3	0.5	0.5
	OXA-24/40 (n=30)	86.7	-	13.3	0.5	4
	OXA-51 (n=30)	100	-	0	0.5	0.5
OXA-58 (n=30)	100	-	0	0.5	1	
Gentamicin ( $\leq 4$   8   $\geq 16$ )	All (n=137)	29.9	3.6	66.4	>32	>32
	MPM <sup>S</sup> (n=17)	82.4	0	17.6	0.5	16
	OXA-23 (n=30)	30	0	70	>32	>32
	OXA-24/40 (n=30)	30	10	60	>32	>32
	OXA-51 (n=30)	3.3	3.3	93.3	>32	>32
OXA-58 (n=30)	26.7	3.3	70	16	>32	
Minocycline ( $\leq 4$   8   $\geq 16$ )	All (n=137)	75.9	10.9	13.1	1	16
	MPM <sup>S</sup> (n=17)	88.2	11.8	0	0.125	8
	OXA-23 (n=30)	50	13.3	36.7	4	16
	OXA-24/40 (n=30)	86.7	10	3.3	1	8
	OXA-51 (n=30)	83.3	3.3	13.3	1	16
OXA-58 (n=30)	76.7	16.7	6.7	0.5	8	
Tigecycline ( $\leq 2$   4   $\geq 8$ )	All (n=137)	59.9	30.7	9.5	2	4
	MPM <sup>S</sup> (n=17)	100	0	0		