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# Introduction

- PLG0206 is a novel engineered cationic antimicrobial peptide being evaluated for treatment of prosthetic joint infections.
- PLG0206 is a broad-spectrum antimicrobial agent with activity against multidrug resistant organisms, has rapid activity against biofilms, and does not have significant local or systemic toxicity in animal models (1,2).
- In this study, the activity of PLG0206 was evaluated by broth microdilution against 104 isolates of Staphylococcus epidermidis, 53 other coagulase-negative staphylococci (CoNS), and 66 Gram-negative isolates consisting of Enterobacterales, Pseudomonas aeruginosa, and Acinetobacter baumannii.

# Methods

- 223 test organisms consisted of reference strains from the American Type Culture Collection, the Centers for Disease Control Antibiotic Reference Bank and clinical isolates from the Micromyx repository.
- Imipenem, levofloxacin, tigecycline, linezolid, vancomycin, oxacillin, ceftazidime, colistin, and amikacin were tested as comparators.
- Testing was conducted in accordance with guidelines from the Clinical and Laboratory Standards Institute (3,4).
- The media employed for testing in the broth microdilution MIC assay for all organisms were cation-adjusted Mueller Hinton Broth (CAMHB) and for PLG0206 only included RPMI-1640 medium supplemented with 0.002% P-80 due to precipitation/flocculation of PLG0206 observed in CAMHB.

# **Methods**

- A standardized inoculum of each organism was prepared per CLSI methods (3, 4).
- Suspensions were prepared in PBS to equal a turbidity of a 0.5 McFarland standard that yielded a final cell density in the plates of approximately 5 x 10<sup>5</sup> CFU/mL.

### Results

Activity of PLG0206 and comparators against CoNS and resistant Gram-negative pathogens are shown in the Table.

Organism	Drug	MIC Range			Mode	MIC <sub>50</sub>	MIC <sub>90</sub>	%S	%R
All S. epidermidis (104)	PLG0206 CAMHB	0.25	-	4	1	1	1	NA	NA
	PLG0206 RPMI	0.03	-	0.12	0.12	0.12	0.12	NA	NA
	Imipenem	< 0.008	-	>8	0.015	0.06	>8	44.2%	55.8%
	Levofloxacin	0.12	-	>4	>4	4	>4	47.1%	52.9%
	Tigecycline	0.12	-	2	0.25	0.25	1	NA	NA
	Vancomycin	1	-	4	2	2	2	100%	0%
	Linezolid	0.5	-	>8	2	2	2	98.1%	1.9%
	Oxacillin	0.06	_	>16	0.12	1	>16	44.2%	55.8%
MSSE (46)	PLG0206 CAMHB	0.25	-	4	1	1	1	NA	NA
	PLG0206 RPMI	0.03	-	0.12	0.12	0.12	0.12	NA	NA
	Imipenem	< 0.008	-	0.06	0.015	0.015	0.015	100%	0%
	Levofloxacin	0.12	-	>4	0.25	0.25	>4	84.8%	15.2%
	Tigecycline	0.12	-	2	0.25	0.25	1	NA	NA
	Vancomycin	1	_	4	2	2	2	100%	0%
	Linezolid	0.5	_	4	2	2	2	100%	0%
	Oxacillin	0.06	-	0.25	0.12	0.12	0.25	100%	0%
MRSE (58) <sup>1</sup>	PLG0206 CAMHB	0.25	-	4	1	1	1	NA	NA
	PLG0206 RPMI	0.03	-	0.12	0.12	0.12	0.12	NA	NA
	Imipenem	< 0.008	-	>8	>8	2	>8	0%	100%
	Levofloxacin	0.12	_	>4	>4	>4	>4	17.2%	82.8%
	Tigecycline	0.12	-	1	1	0.5	1	NA	NA
	Vancomycin	1	-	4	2	2	2	100%	0%
	Linezolid	1	-	>8	2	2	4	96.6%	3.4%
	Oxacillin	0.5	-	>16	>16	16	>16	0%	100%
CoNS non- epidermidis (53) <sup>2</sup>	PLG0206 CAMHB	< 0.12	-	4	0.25	0.25	1	NA	NA
	PLG0206 RPMI	0.015	-	2	0.06	0.06	0.06	NA	NA
	Imipenem	< 0.008	_	>8	0.03	0.03	0.25	52.8%	47.2%
	Levofloxacin	0.12	-	>4	0.5	0.5	>4	83.0%	17%
	Tigecycline	0.25	-	2	0.5	0.5	1	NA	NA
	Vancomycin	0.5	-	2	1	1	1	100%	0%
	Linezolid	1	-	4	2	2	4	100%	0%
	Oxacillin	0.12	_	>16	0.12	0.5	8	52.8%	47.2%

<u>Results</u>										
Organism	Drug	MIC Range			Mode	MIC <sub>50</sub>	MIC <sub>90</sub>	%S	%R	
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Enterobacterales (22) <sup>3</sup>	PLG0206 CAMHB	1	-	>8	8	8	>8	NA	NA	
	PLG0206 RPMI	< 0.12	_	>128	0.5	0.5	32	NA	NA	
	Imipenem	2	-	>8	>8	>8	>8	0%	95.5%	
	Levofloxacin	0.06	-	>4	>4	>4	>4	14%	86%	
	Tigecycline	0.25	-	4	0.5	0.5	2	95.4%	0%	
	Ceftazidime	0.12	-	>32	>32	>32	>32	5%	86%	
	Colistin	0.06	-	>16	0.25	0.12	16	NA	NA	
	Amikacin	1	-	>64	>64	16	>64	55%	32%	
P. aeruginosa (20)	PLG0206 CAMHB	8	-	>8	8	8	>8	NA	NA	
	PLG0206 RPMI	0.5	-	4	1	1	2	NA	NA	
	Imipenem	4	-	>8	>8	>8	>8	0%	90%	
	Levofloxacin	0.5	-	>4	>4	>4	>4	10%	90%	
	Tigecycline	8	_	>16	>16	>16	>16	NA	NA	
	Ceftazidime	4	-	>32	>32	>32	>32	10%	85%	
	Colistin	0.12	_	2	0.5	0.5	1	100%	0%	
	Amikacin	0.5	_	>64	8	8	>64	55%	30%	
A. baumannii (24)	PLG0206 CAMHB	2	_	8	4	4	8	NA	NA	
	PLG0206 RPMI	0.25	-	0.5	0.25	0.25	0.5	NA	NA	
	Imipenem	1	-	>8	>8	>8	>8	8.3%	87.5%	
	Levofloxacin	4	-	>4	>4	>4	>4	0%	96%	
	Tigecycline	1	-	16	4	4	4	NA	NA	
	Ceftazidime	>32	-	>32	>32	>32	>32	0%	100%	
	Colistin	0.12	-	>16	0.25	0.25	4	83%	17%	
	Amikacin	2	_	>64	>64	>64	>64	21%	79%	

negative staphylococci. NA, not applicable. %S, percentage of isolates that are susceptible according to established CLSI breakpoints (tigecycline FDA breakpoints used where applicable). %R, percentage of isolates that are resistant according to established CLSI breakpoints (tigecycline FDA breakpoints used where applicable).

For testing PLG0206 in RPMI one isolate did not grow; data for PGL0206 RPMI is based on 57 isolates

Species breakdown for non-epidermidis CoNS- S. hominis (10), S. haemolyticus (10), S. warneri (8), S. capitis (4), S. simulans (5), S. lugdunensis (4), S. caprae (4), S. saprophyticus (7), and S. pettenkoferi (1).

Species breakdown for Enterobacterales E. coli (8), K. pneumoniae (8), P. mirabilis (2), E. cloacae (2), K. oxytoca (1), C. freundii (1).

- Overall, the PLG0206 MIC values were much lower when the organisms were tested in RPMI relative to CAMHB. For example, the MIC50 and MIC90 values were 4 to 16-fold lower in RPMI than CAMHB for the tested groups of organisms (CoNS, Enterobacterales, *P. aeruginosa*, and *A. baumannii*) **(Table 1).**
- Methicillin-resistance in *S. epidermidis* did not affect the MIC50/MIC90 values or MIC distribution of PLG0206.

### **Conclusions**

- •PLG0206 was found to have potent antimicrobial activity against CoNS; activity in RPMI was more potent than CAMHB. There was no impact of methicillin-resistance among CoNS on the activity of PLG0206.
- •PLG0206 had varying activity against resistant Gram-negative bacilli. Among Enterobacterales, PLG0206 was most active against E. coli, E. cloacae, and C. freundii, was comparatively less active against *Klebsiella* spp., and was inactive against *P. mirabilis*.
- •Despite the high degree of drug resistance among the evaluated *P*. aeruginosa and A. baumannii, PLG0206 had potent activity when evaluated in RPMI. As with CoNS, PLG0206 MIC values were several-fold higher in CAMHB relative to RPMI when testing Gram-negative bacilli.

#### References

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