

05. New antibacterial agents, PK/PD & Stewardship  
5a. Mechanisms of action, new compounds, preclinical data & pharmacology of antibacterial agents

# Resistance development to PLG0206, a novel engineered antimicrobial peptide

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## Background

- PLG0206 is an engineered antimicrobial peptide (1, 2) with rapid bactericidal, broad-spectrum, and anti-biofilm properties being developed for the treatment of periprosthetic joint infections (PJI). The aim of this study was to determine the spontaneous mutation frequency (SMF) of PLG0206 with a variety of Gram-positive and Gram-negative pathogens that can cause PJs.

## Methods

- Isolates (*S. aureus*, *S. epidermidis*, enterococci, *S. pyogenes*, and *P. aeruginosa*) were sourced from the ATCC and Micromyx strain repository.
- Bacteria were plated onto RPMI-1640 agarose containing PLG0206 at 4, 8, or 16-fold the agar dilution MIC value.
- After a 48 hour incubation, the SMF was calculated and putative spontaneous mutant colonies were picked and patched to fresh agar plates containing the same PLG0206 concentration.
- Putative spontaneous mutant broth microdilution MIC testing utilized RPMI for PLG0206 while comparators were tested in CLSI standard media (3, 4).

## Results

Table 1. Agar dilution MIC of isolates

Organism	Isolate	Type	MIC (µg/ml)				
			PLG0206 <sup>1</sup>	GM	IPM	LVX	VAN
<i>S. aureus</i>	ATCC 29213	MSSA, QC	1	0.25 (0.12-1) <sup>2</sup>	0.016 (0.016-0.06) <sup>2</sup>	0.25 (0.06-0.5) <sup>2</sup>	1 (0.5-2) <sup>2</sup>
	NRS130	MSSA; ERY <sup>R</sup>	2	0.25	0.008	0.25	1
	NRS169	MSSA; ERY <sup>R</sup> , CLI <sup>R</sup>	1	0.25	0.016	0.12	1
	NRS1/Mu50	MRSA; VISA	2	>4	>8	8	4
	NRS384 (USA300)	MRSA	2	0.25	0.25	0.5	1
	NRS382 (USA100)	MRSA	1	0.25	1	4	1
<i>S. epidermidis</i>	MMX 8740	MRSE	1	8	2	16	2
	MMX 8655	MRSE	1	0.25	0.016	4	2
ATCC 49134	MSSE		0.5	0.06	0.008	0.25	2
<i>E. faecalis</i>	ATCC 700802	V583 VRE		<2.94E-10	<2.94E-10	<2.94E-10	<4
	MMX 0485	VanA VRE		<8.93E-10	<8.93E-10	<8.93E-10	<4
<i>S. pyogenes</i>	ATCC 700294	-		<4.55E-10	<4.55E-10	<4.55E-10	<128
	ATCC 27853	QC		8.00E-08	6.81E-08	1.27E-08	>32
<i>P. aeruginosa</i>	CDC 0241	IMP-1		8.67E-08	1.71E-07	1.13E-07	>32
	MMX 3025	CIP <sup>R</sup>		6.80E-08	5.33E-09	<8.20E-10	64

## Results

Table 2. Spontaneous mutation frequency values

Organism	Isolate	Type	Spontaneous Mutation Frequency at X-fold the MIC			MPC (µg/ml)
			4x	8x	16x	
<i>S. aureus</i>	ATCC 29213	MSSA; QC	2.01E-07	4.54E-08	1.03E-09	>16
	NRS130	MSSA; ERY <sup>R</sup>	1.76E-08	1.18E-09	<1.18E-09	32
	NRS169	MSSA; ERY <sup>R</sup> , CLI <sup>R</sup>	3.14E-08	4.24E-09	<8.47E-10	16
	NRS1	MRSA; VISA	<1.98E-09	<1.98E-09	<1.98E-09	<8
	NRS384	MRSA (USA300)	3.48E-08	7.09E-10	<7.09E-10	32
	NRS382	MRSA (USA100)	2.39E-08	2.84E-08	<1.14E-09	16
<i>S. epidermidis</i>	MMX 8740	MRSE	NA	<1.22E-09	<1.22E-09	8
	MMX 8655	MRSE	NA	<1.19E-09	<1.19E-09	8
	ATCC 49134	MSSE	NA	NA	<1.19E-09	8
<i>E. faecalis</i>	ATCC 700802	V583 VRE		<2.94E-10	<2.94E-10	<2.94E-10
<i>E. faecium</i>	MMX 0485	VanA VRE		<8.93E-10	<8.93E-10	<8.93E-10
<i>S. pyogenes</i>	ATCC 700294	-		<4.55E-10	<4.55E-10	<4.55E-10
<i>P. aeruginosa</i>	ATCC 27853	QC		8.00E-08	6.81E-08	1.27E-08
	CDC 0241	IMP-1		8.67E-08	1.71E-07	1.13E-07
	MMX 3025	CIP <sup>R</sup>		6.80E-08	5.33E-09	<8.20E-10

Table 3. Broth dilution MIC values for putative *S. aureus* spontaneous mutants

Isolate	Selection Concentration at X-fold the MIC	Isolate Name	PLG0206 <sup>1</sup>	VAN	RIF	ERY	CLI	LZD	LVX	TET
				N/A	4X	8X	16X			
ATCC 29213	N/A	parent	0.5	1	0.008	0.5	0.25	4	0.25	1
	4X	P-4-100-A	1	1	0.008	0.25	0.25	4	0.12	0.5
	4X	P-4-100-B	1	0.5	0.008	0.25	0.25	4	0.12	0.5
	8X	P-8-100-A	1	0.5	0.008	0.25	0.25	4	0.12	0.5
	8X	P-8-100-B	1	0.5	0.008	0.25	0.25	4	0.12	0.5
	16X	P-16-100-1	1	0.5	0.008	0.25	0.12	4	0.12	0.5
NRS130	N/A	parent	0.5	0.5	0.008	>32	0.12	2	0.12	0.25
	4X	P-8-2131-A	1	0.5	0.008	>32	0.25	2	0.12	0.25
	4X	P-8-2131-B	1	1	0.008	>32	0.25	2	0.25	0.25
NRS169	N/A	parent	0.5	0.5	0.016	0.25	0.12	4	0.12	0.25
	8X	P-8-2134-A	0.5	1	0.016	0.5	0.25	4	0.12	0.25
	8X	P-8-2134-B	0.5	1	0.016	0.5	0.25	4	0.12	0.25
NRS384/ MMX 2011	N/A	parent	0.5	0.5	0.008	>32	0.12	2	0.5	32
	4X	P-8-2011-A	0.5	0.5	0.008	>32	0.12	2	0.5	32
	8X	P-16-2011-1	0.5	1	0.008	>32	0.12	2	1	32
NRS382/ MMX 2119	N/A	parent	0.5	1	0.016	>32	0.12	2	0.5	32
	4X	P-4-2119-A	0.5	1	0.008	>32	>4	2	4	0.25
	8X	P-8-2119-A	1	1	0.016	>32	>4	2	4	0.25

<sup>1</sup> MIC as determined in RPMI-1640 Agar; <sup>2</sup> CLSI QC ranges shown in parentheses where applicable; GM: gentamicin, IPM: imipenem, LVX: levofloxacin, VAN: vancomycin, MSSA: methicillin-susceptible *S. aureus*, MRSA: methicillin-resistant *S. aureus*, VISA: vancomycin-intermediate *S. aureus*, QC: quality control CIP<sup>R</sup>: ciprofloxacin-resistant, ERY<sup>R</sup>: erythromycin-resistant, CLI<sup>R</sup>: clindamycin-resistant

## Results

Table 4. Broth dilution MIC values for *P. aeruginosa* spontaneous mutants

Isolate	Selection Concentration at X-fold the MIC	Isolate Name	PLG0206<sup>1</sup>	IMP	COL	AZT	LVX	GM	CAZ	TZP<sup>2</sup>
N/A	8-4X	16-8X	32-16X							




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