## INTRODUCTION

- PLG0206 is an investigational, engineered cationic antimicrobial peptide designed to overcome the shortcomings of other natural AMPs, such as toxicity and limited activity (1, 2). PLG0206 has recently been shown to be well tolerated and safe in a phase 1 study (3).
- This study evaluated the activity of PLG0206 and comparator antimicrobials against non-fermenting Gram-negative rods from the IHMA repository of isolates collected from various world-wide locations in 2019.

## MATERIALS AND METHODS

- Isolates tested included Acinetobacter baumannii (298), Pseudomonas aeruginosa (300) and Stenotrophomonas *maltophilia* (150). The isolates were collected in 2019, with approximately one-third from Europe and the remainder from other regions (Figure 1). The isolates originated mainly from respiratory infections, but also included a variety of other infection types (Figure 2).
- Minimum inhibitory concentrations (MICs) were determined by Clinical and Laboratory Standards Institute (CLSI) broth microdilution methodology (4) in cation-adjusted Mueller Hinton broth (CA-MHB), except for PLG0206 which was tested in MOPS RPMI-1640 medium supplemented with 0.002% Tween-80 due to precipitation of PLG0206 observed in CAMHB.
- The susceptibility of comparators was determined using the 2022 CLSI breakpoints (5).
- Multi-drug-resistance (MDR) for A. baumannii and P. aeruginosa was defined as resistance to 3 or more of the antimicrobials tested by class of antimicrobial including aminoglycosides (amikacin, gentamicin or tobramycin), cephems (cefepime or ceftazidime), fluoroquinolones (levofloxacin or ciprofloxacin) and tetracyclines (doxycycline or tetracycline) or by individual antimicrobial for aztreonam, ampicillin/sulbactam, colistin, ceftazidime/avibactam, meropenem and piperacillin/tazobactam. MDR for *S. maltophilia* was defined as resistance to both ceftazidime and levofloxacin (the only agents tested with CLSI) breakpoints).

#### FIGURE 1: Geographical location of the non-fermenting Gram-negative rods tested.

# rods

mikacin

treonam

listin xycyclin

PLG0206

Amikacin

μg/ml) Cefepime Ceftazidin

Colistin

Doxycycl entamici

(8 µg/ml) Piperacilli

μg/ml) Tetracyclin



FIGURE 2: Infection source for non-fermenting Gram-negative rods tested.



#### FIGURE 3: Cumulative % MIC distribution for PLG0206 against non-fermenting Gram-negative rods



#### TABLE 1: Summary of activity of PLG0206 and comparators against non-fermenting Gram-negative

		A. baun	nannii (	n=298)			P. aerug	jinosa (	n=297)			S. malto	philia (	n=89)		
	MIC <sub>50</sub>	MIC <sub>90</sub>	%S	%I	%R	MIC <sub>50</sub>	MIC <sub>90</sub>	%S	%I	%R	MIC <sub>50</sub>	MIC <sub>90</sub>	%S	%I	%R	
	0.5	1	-	-	-	1	2	-	-	-	0.5	1	-	-	-	
	>64	>64	39.9	2.3	57.7	4	>64	85.2	2.4	12.5	>64	>64	-	-	-	
actam (2:1)	32	>32	66.4		N	ot tested	ł		Not tested							
		N		8	>16	69.0	11.1	19.9	Not tested							
oactam (4 µg/ml)	Not tested					4	>16	-	-	-	Not tested					
	>32	>32	26.2	2.3	71.5	4	>32	75.4	7.1	17.5	>32	>32	-	-	-	
	>32	>32	29.5	1.7	68.8	4	>32	71.4	5.7	22.9	>32	>32	33.7	3.4	62.9	
bactam (4 µg/ml)		N	ot tested	b		2	>16	87.2	-	12.8	Not tested					
	>32	>32	27.2	0.3	72.5	0.12	16	76.1	4.0	19.9	4	16	-	-	-	
	0.25	0.5	-	97.3	2.7	0.5	1	-	99.7	0.3	8	>32	-	-	-	
	4	64	57.0	5.0	37.9	8	32	-	-	-	2	4	-	-	-	
	>32	>32	38.3	3.4	58.4	1	>32	82.5	3.0	14.5	>32	>32	-	-	-	
	16	>32	28.2	5.4	66.4	0.5	32	69.4	9.1	21.5	2	8	79.8	7.9	12.4	
	64	>64	29.9	1.3	68.8	0.5	32	77.4	7.1	15.5	>64	>64	-	-	-	
oorbactam (8 µg/ml)	>32	>32	-	-	-	0.5	32	-	-	-	>32	>32	-	-	-	
obactam (4 µg/ml)	>128	>128	26.8	3.0	70.1	8	128	73.4	12.1	14.5	>128	>128	-	-	-	
	32	>32	33.6	11.1	55.4	8	32	-	-	-	16	32	-	-	-	
	16	>32	45.3	2.0	52.7	0.5	>32	84.5	0.7	14.8	>32	>32	-	-	-	

TABLE 2: Summary of activity of PLG0206 and comparators against multi-drug-resistant (MDR) nonfermenting Gram-negative rods.

MRC5         MRC9 $\odot$ $\checkmark$ $\checkmark$ $MRC5$ $MRC9$ $\circ$ $\checkmark$ $MRC9$ $\circ$ $MRC90$ $\circ$ $\sim$ $\sim$ $\circ$ $\circ$ $\sim$ $\circ$		<i>A. baumannii</i> (n=216)						P. aerug	jinosa	(n=70	)	S. maltophilia (n=10)					
0.51110.510.510.5100264>6417.13.279.664>6426442.95.751.4>64>64>6411026751.4>64>64>64-1-1-236361261361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361361	l	MIC <sub>5</sub>	0 MIC	%S	%I	%R	0 MIC	0 MIC	%S	%I	%R	MIC <sub>50</sub>	MIC <sub>90</sub>	% S	%I	%R	
>64       >64       17.1       3.2       79.6       64       >64       42.9       5.7       51.4       >64       >64       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       - <th< th=""><th></th><th>0.5</th><th>1</th><th>-</th><th>-</th><th>-</th><th>1</th><th>1</th><th>-</th><th>-</th><th>-</th><th>0.5</th><th>1</th><th>-</th><th>-</th><th>-</th></th<>		0.5	1	-	-	-	1	1	-	-	-	0.5	1	-	-	-	
Ibactam (2:)       >32       >32       5.6       91.2 $ispace       ispace       $		>64	>64	17.1	3.2	79.6	64	>64	42.9	5.7	51.4	>64	>64	-	-	-	
Vibactam (4	Ibactam (2:1)	>32	>32	3.2	5.6	91.2		No	ot teste	d		Not tested					
vibactam (4       Not tested       16       >16       >16       -       -       -       -       Not tested       -       -       Not tested       -       -       Not tested       -       -       Not tested       -       -       -       Not tested       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -       -		Not tested					>16	>16	14.3	17.1	68.6	Not tested					
>32       >32       0.5       2.3       97.2       32       >32       1.4       21.4       67.1       >32       >32            >32       >32       3.7       1.4       94.9       >32       >32       10.0       10.0       80.0       >32       >32       0.0       0.0       100.0         avibactam(4       Image:	vibactam (4		No	d		16	>16	-	-	-	Not tested						
>32       >32       3.7       1.4       94.9       >32       >32       10.0       10.0       80.0       >32       >32       0.0       10.0         avibactam (4       Vortester       16       >16       45.7       1.4       64.3       16       >32       >32       0.0       0.0       100.0         avibactam (4          16       >32       28.6       7.1       64.3       16       >32              96.8       3.2       1       1        98.6       1.4       16       >32               96.8       3.2       1       1        98.6       1.4       16       >32		>32	>32	0.5	2.3	97.2	32	>32	11.4	21.4	67.1	>32	>32	-	-	-	
avibactam (4)       Image: Substrate		>32	>32	3.7	1.4	94.9	>32	>32	10.0	10.0	80.0	>32	>32	0.0	0.0	100.0	
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166441.76.951.416>128416>32>3215.74.280.132>3240.02.957.1>32>32>3216>322.87.489.816>3230.07.162.98>320.00.0100.0200>64>643.21.994.916>6428.68.662.9>64>64200 $322$ $332$ $1.9$ 94.916>6428.68.662.9>64>64200 $332$ $332$ $1.9$ 94.916>6428.68.662.9>64>64200 $332$ $332$ $1.9$ 94.916>6428.68.662.9>64>64200 $332$ $332$ $1.9$ 94.916>32 $322$ $1.2$ $3.2$ $332$ $2.32$ $1.0$ 0.0100.0200 $332$ $332$ $1.2$ $1.2$ $1.2$ $2.3$ $3.2$ $2.32$ $2.32$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ $1.2$ <th></th> <th>0.25</th> <th>0.5</th> <th>-</th> <th>96.8</th> <th>3.2</th> <th>1</th> <th>1</th> <th>-</th> <th>98.6</th> <th>1.4</th> <th>16</th> <th>&gt;32</th> <th>-</th> <th>-</th> <th>-</th>		0.25	0.5	-	96.8	3.2	1	1	-	98.6	1.4	16	>32	-	-	-	
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azobactam (4)       >128       >128       0.9       2.3       96.8       128       >12.8       12.9       30.0       57.1       >128       >128       -       -       -         >32       >32       >32       11.6       13.9       74.5       0.5       1       -       -       0.5       0.5       0.5       -       -         >32       >32       232       24. 5       2.8       72.7       32       >32       38. 6       2.9       58.6       >32       >32       -       -       -	vaborbactam	>32	>32	-	-	-	16	>32	-	-	-	>32	>32	-	-	-	
>32       >32       11.6       13.9       74.5       0.5       1       -       -       0.5       0.5       -       -       -         >32       >32       24.       2.8       72.7       32       >32       38.       2.9       58.6       >32       >32       -       -       -	azobactam (4	>128	>128	0.9	2.3	96.8	128	>128	12.9	30.0	57.1	>128	>128	-	-	-	
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		>32	>32	24. 5	2.8	72.7	32	>32	38. 6	2.9	58.6	>32	>32	-	-	-	

### **RESULTS SUMMARY**

• 3 *P. aeruginosa* and 61 *S. maltophilia* were unable to grow in RPMI, so all antimicrobial data presented exclude these isolates. All A. baumannii remained viable in RPMI. • PLG0206 was very active against *A. baumannii*, *S. maltophilia,* and *P. aeruginosa* (Figure 3 & Table 1). • PLG0206 retained activity against MDR isolates of A. baumannii, P. aeruginosa and S. maltophilia (Table 2).

### CONCLUSIONS

### References

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• PLG0206 was the most active antimicrobial overall (based on MIC<sub>50</sub> and MIC<sub>90</sub>) against *S. maltophilia* and was the most active overall against A. baumannii and P. aeruginosa except for colistin. However, the use of colistin is not ideal due to its safety and tolerability profile, and recent

breakpoint changes render all isolates to be of intermediate susceptibility to colistin at best.

• As a high proportion of *S. maltophilia* were unable to grow in RPMI, further work is underway to find an alternative test medium for PLG0206.

 These data suggest that this novel antimicrobial peptide could be a potential treatment option for these difficult to treat bacterial pathogens, including MDR strains.