

# Supplemental Table 1

Supplemental Table 1. *P. aeruginosa* strain isolate information.

<b><i>P. aeruginosa</i> Strain</b>	<b>Source</b>
<b>PAO1</b>	Infected wound (ATCC 15692)
<b>PA103</b>	Sputum (ATCC 29260)
<b>H25815</b>	Tracheal aspirate
<b>H28822</b>	Catheterized urine
<b>M57-15</b>	Mucoid CF isolate
<b>M69781</b>	Sputum
<b>T63621</b>	Tracheal aspirate
<b>W40423</b>	Sputum
<b>W41033</b>	Urine
<b>W43532</b>	Bronchial washing

## Supplemental Table 2

Target	NCBI Gene ID	Number Designation	Base Sequence (5' to 3')	Location on gene relative to start site	5' End	3' End
<i>AcpP</i>	879895	-0206	CAT ACC TTG TT	-8 to +3	TEG	(RXR) <sub>4</sub> XB
<i>AcpP</i>	879895	-0307	CTC ATA CCT TG	-6 to +5	H	(RXR) <sub>4</sub> XB
<i>AcpP</i>	879895	-0445	CTC ATA CCT TG	-6 to +5	(RXR) <sub>4</sub> XB	H
<i>AcpP</i>	879895	-0446	CTC ATA CCT TG	-6 to +5	(RGR) <sub>4</sub> XB	H
<i>AcpP</i>	879895	-0447	CTC ATA CCT TG	-6 to +5	(RFR) <sub>4</sub> XB	H
<i>LpxC</i>	881292	-0251	GTT GTT TGA TC	+3 to +13	(RXR) <sub>4</sub> XB	H
<i>LpxC</i>	881292	-0505	GTT GTT TGA TC	+3 to +13	TEG	(RXR) <sub>4</sub> XB
<i>RpsJ</i>	881717	-0066	CCT CAG ACT CC	-15 to -5	(RXR) <sub>4</sub> XB	H
<i>RpsJ</i>	881717	-0067	CCT CAG ACT CC	-15 to -5	R6G	H
<i>RpsJ</i>	881717	-0207	CCT CAG ACT CC	-15 to -5	TEG	(RGR) <sub>4</sub> XB
<i>RpsJ</i>	881717	-0208	CCT CAG ACT CC	-15 to -5	TEG	R6G
<i>RpsJ</i>	881717	-0209	GCA TTT GAC CT	-7 to +4	TEG	(RXR) <sub>4</sub> XB
<i>RpsJ</i>	881717	-0249	GCA TTT GAC CT	-7 to +4	(RXR) <sub>4</sub> XB	H
<i>RpsJ</i>	881717	-0308	CCT CAG ACT CC	-15 to -5	H	(RXR) <sub>4</sub> XB
<i>RpsJ</i>	881717	-0371	GCA TTT GAC CT	-7 to +4	R6G	H
<i>RpsJ</i>	881717	-0628	CCT CAG ACT CC	-15 to -5	(dRdFdF) <sub>3</sub> XB (D amino acids)	Acetyl
<i>RpsJ</i>	881717	-0629	CCT CAG ACT CC	-15 to -5	(dRdFdF) <sub>3</sub> dRXB (D amino acids)	Acetyl
<i>RpsJ</i>	881717	-0630	CCT CAG ACT CC	-15 to -5	(dRXdR) <sub>4</sub> XB	Acetyl
<i>RpsJ</i>	881717	-0631	CCT CAG ACT CC	-15 to -5	R8B (D amino acids)	Acetyl
<i>RpsJ</i>	881717	-0632	CCT CAG ACT CC	-15 to -5	dR6G (D amino acids)	Acetyl
<i>RpsJ</i>	881717	-1075	CCT CAG ACT CC	-15 to -5	TEG	H
<i>Scr</i>	N/A	-0078	ATC GTT GCA TC	N/A	(RXR) <sub>4</sub> XB	H
<i>Scr</i>	N/A	-0949	TCT CAG ATG GT	N/A	TEG	(RXR) <sub>4</sub> XB
<i>Scr</i>	N/A	-1073	TCT CAG ATG GT	N/A	(RXR) <sub>4</sub> XB	H

### Supplemental Table 2. PPMOs.

PPMOs are listed in alphabetical and numerical order. For the location relative to the start site we defined 'A' of ATG as +1. Non-common abbreviations are as follows – X: 6-amino hexanoic acid (aminocaproic acid), B: Beta-alanine (for conjugation), d: prefix for non-natural stereoisomer form of an amino acid, TEG: Triethylene glycol, Scr: Scramble.

# Supplemental Figure 1

(a) MICs of PPMOs in MH II media

Strain ID	Species	PPMOs												
		AcpP-0206	AcpP-0445	LpxC-0251	LpxC-0505	RpsJ-0066	RpsJ-0207	RpsJ-0208	RpsJ-0209	RpsJ-0628	RpsJ-0629	RpsJ-0630	RpsJ-0632	
PAO1	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
H25815	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
H28822	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
M57-15	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
M69781	<i>P. aeruginosa</i>	>16	NT	NT	8	NT	4	>16	>16	>16	>16	>16	>16	>16
T63621	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
W40423	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
W41033	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
W43532	<i>P. aeruginosa</i>	>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16
PPMO MIC <sub>50</sub> [μM]		>16	NT	NT	>16	NT	>16	>16	>16	>16	>16	>16	>16	>16

Key

NT	Not Tested
>16	>16 μM
16	16 μM
8	8 μM
4	4 μM
2	2 μM
1	1 μM
≤0.5	≤0.5 μM

(b) MICs of PPMOs in MH II media with 1μg/ml Colistin

Strain ID	Species	PPMOs												Colistin MIC (μg/mL)	
		AcpP-0206	AcpP-0445	LpxC-0251	LpxC-0505	RpsJ-0066	RpsJ-0207	RpsJ-0208	RpsJ-0209	RpsJ-0628	RpsJ-0629	RpsJ-0630	RpsJ-0632		
PAO1	<i>P. aeruginosa</i>	8	NT	NT	2	NT	8	>16	>16	8	>16	2	8	4	4
H25815	<i>P. aeruginosa</i>	≤0.5	NT	NT	≤0.5	NT	≤0.5	16	16	≤0.5	≤0.5	≤0.5	≤0.5	≤0.5	8
H28822	<i>P. aeruginosa</i>	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	NT	2
M57-15	<i>P. aeruginosa</i>	1	NT	NT	≤0.5	NT	>16	4	≤0.5	>16	>16	>16	>16	1	4
M69781	<i>P. aeruginosa</i>	≤0.5	NT	NT	≤0.5	NT	>16	1	≤0.5	≤0.5	≤0.5	≤0.5	≤0.5	≤0.5	8
T63621	<i>P. aeruginosa</i>	4	NT	NT	2	NT	>16	8	2	4	8	2	2	2	2
W40423	<i>P. aeruginosa</i>	4	NT	NT	2	NT	>16	16	2	8	16	2	2	4	8
W41033	<i>P. aeruginosa</i>	≤0.5	NT	NT	≤0.5	NT	16	2	≤0.5	≤0.5	8	≤0.5	≤0.5	1	4
W43532	<i>P. aeruginosa</i>	2	NT	NT	≤0.5	NT	≤0.5	≤0.5	≤0.5	>16	>16	>16	>16	2	4
PPMO MIC <sub>50</sub> [μM]		1	NT	NT	≤0.5	NT	16	4	≤0.5	4	8	2	2	1	4

(c) MICs of PPMOs in MH II media with 2μg/ml PMBN

Strain ID	Species	PPMOs												PMBN MIC (μg/mL)	
		AcpP-0206	AcpP-0445	LpxC-0251	LpxC-0505	RpsJ-0066	RpsJ-0207	RpsJ-0208	RpsJ-0209	RpsJ-0628	RpsJ-0629	RpsJ-0630	RpsJ-0632		
PAO1	<i>P. aeruginosa</i>	4	2	2	2	2	>16	>16	4	>16	>16	4	8	>16	>16
H25815	<i>P. aeruginosa</i>	≤0.25	≤0.5	1	1	≤0.5	≤0.5	1	≤0.5	2	4	≤0.5	1	1	>16
H28822	<i>P. aeruginosa</i>	16	2	2	1	8	>16	>16	4	>16	>16	4	16	>16	>16
M57-15	<i>P. aeruginosa</i>	2	1	2	2	2	>16	>16	16	>16	>16	4	8	>16	>16
M69781	<i>P. aeruginosa</i>	8	1	2	≤0.5	2	1	1	≤0.5	8	16	≤0.5	≤0.5	4	4
T63621	<i>P. aeruginosa</i>	8	1	2	8	2	>16	>16	8	>16	>16	4	8	>16	4
W40423	<i>P. aeruginosa</i>	4	1	8	8	4	>16	>16	16	>16	>16	8	>16	>16	4
W41033	<i>P. aeruginosa</i>	4	1	4	16	2	4	>16	16	>16	>16	8	8	16	>16
W43532	<i>P. aeruginosa</i>	4	1	4	4	2	16	16	16	>16	>16	16	16	>16	>16
PPMO MIC <sub>50</sub> [μM]		4	1	2	2	2	>16	>16	4	>16	>16	4	8	>16	>16

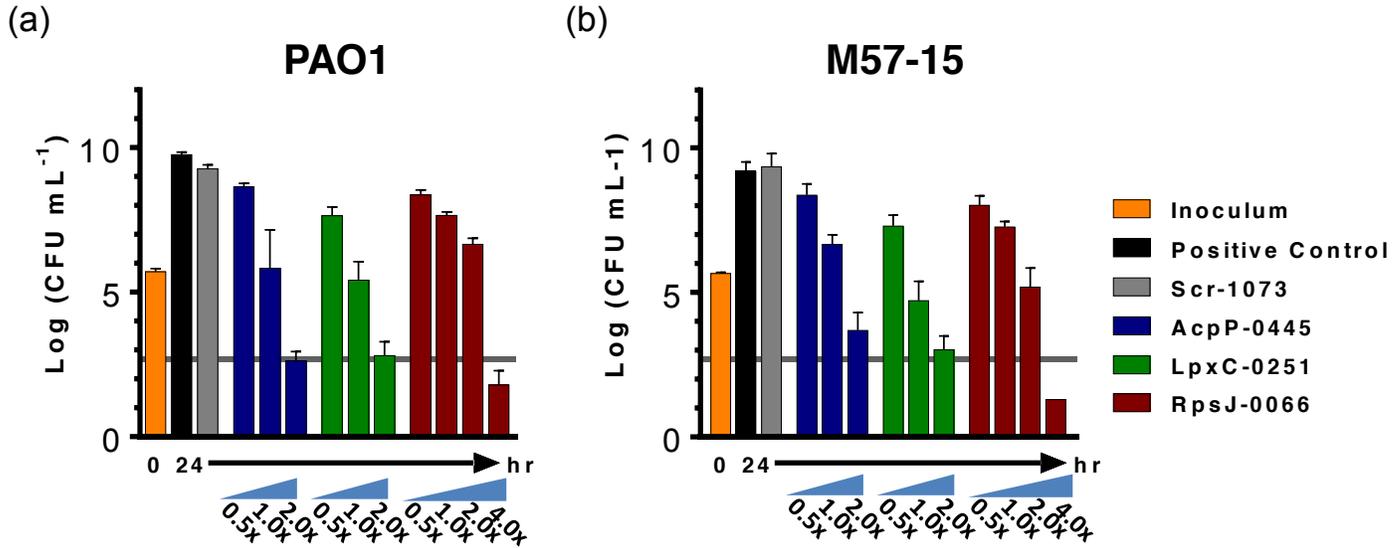
(d) MICs of PPMOs in MOPS media

Strain ID	Species	PPMOs												
		AcpP-0206	AcpP-0445	LpxC-0251	LpxC-0505	RpsJ-0066	RpsJ-0207	RpsJ-0208	RpsJ-0209	RpsJ-0628	RpsJ-0629	RpsJ-0630	RpsJ-0632	
PAO1	<i>P. aeruginosa</i>	1	1	2	4	1	16	>16	8	>16	>16	4	8	>16
H25815	<i>P. aeruginosa</i>	2	2	4	2	1	16	8	8	>16	>16	2	2	4
H28822	<i>P. aeruginosa</i>	4	1	2	4	1	8	>16	4	>16	>16	8	8	>16
M57-15	<i>P. aeruginosa</i>	2	1	2	4	≤0.5	8	>16	8	>16	>16	4	8	16
M69781	<i>P. aeruginosa</i>	4	1	1	2	≤0.5	1	4	1	>16	>16	2	1	4
T63621	<i>P. aeruginosa</i>	1	1	4	4	1	16	>16	4	>16	>16	4	2	>16
W40423	<i>P. aeruginosa</i>	≤0.5	1	1	1	≤0.5	1	2	1	>16	>16	1	2	1
W41033	<i>P. aeruginosa</i>	4	2	4	8	1	8	>16	4	>16	>16	8	2	>16
W43532	<i>P. aeruginosa</i>	2	1	2	8	≤0.5	16	>16	4	>16	>16	8	4	>16
PPMO MIC <sub>50</sub> [μM]		2	1	2	4	1	8	>16	4	>16	>16	4	2	>16

## Supplemental Figure 1. Heatmaps of *P. aeruginosa* growth inhibition by PPMOs.

Heatmaps of PPMO MICs in *P. aeruginosa* strains grown in MHII (a), MHII in the presence of sub-inhibitory concentrations of colistin (b), MHII in the presence of sub-inhibitory concentrations of PMBN (c), or MOPS minimal media (d). Tested PPMOs had minimal activity against *Pseudomonas* grown in MHII alone but the addition of sub-inhibitory concentrations of colistin, PMBN, or growth in MOPS resulted in enhanced antimicrobial inhibition.

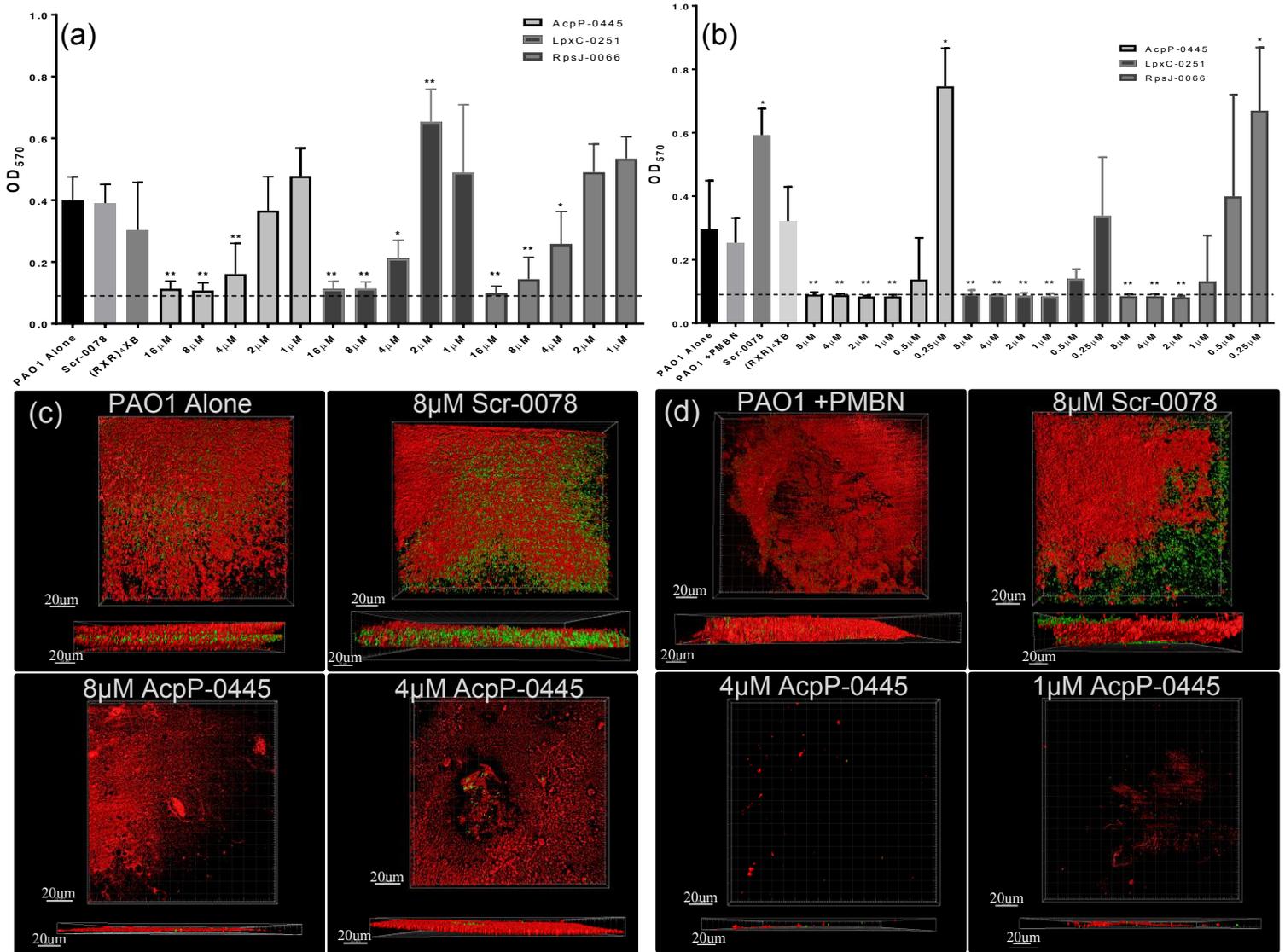
## Supplemental Figure 2



### Supplemental Figure 2: Lead PPMOs against *P. aeruginosa* are bactericidal.

MBC assays of lead PPMOs against PAO1 (a) and M57-15 (b), at 0.5-, 1-, 2-, and 4-times the MIC. Grey-line underlays represent a 3-log CFU reduction as compared to starting inoculum. Positive control is 24 hr culture growth without any treatment. These data are combined results from at least 3 experiments.

# Supplemental Figure 3



## Supplemental Figure 3: PPMO treatment prevents formation of *P. aeruginosa* biofilm.

PAO1-GFP was grown in MBEC plates in the conditions indicated. Optical density measurements of crystal violet stained biofilm formations grown in MHII (a) or in MHII with 2 µg/mL PMBN (b). Dashed line represents the lower limit of detection. This data is combined from 3 independent experiments, error bars represent standard deviation, and statistics were done with a one-way ANOVA and Holm-Šidák's multiple comparisons test where \*\*  $p < 0.0001$  and \*  $p < 0.05$  is indicated for (a) as compared to PAO1 Alone (a) and as compared PAO1+PMBN (b). Spinning disk confocal microscopy images of MBEC pegs are depicted from above and the side incubated in MHII (c), or MHII with 2 µg/mL PMBN (d). Green indicates PAO1-GFP and red is biofilm stained with Concanavalin A-Alexafluor 647.

## Supplemental Figure 4

PPMO ( $\mu$ M)	<i>P. aeruginosa</i> Strain			Antibiotic ( $\mu$ g/mL)	<i>P. aeruginosa</i> Strain		
	PAO1	M57-15	W43532		PAO1	M57-15	W43532
<b>AcpP-0445</b>	8	8	4	<b>Pip/Tazo</b>	4.00	4.00	4.00
<b>LpxC-0251</b>	16	16	8	<b>Tobramycin</b>	0.50	0.50	0.50
<b>RpsJ-0066</b>	8	32	32	<b>Levofloxacin</b>	0.50	0.25	0.25
<b>Scr-1073</b>	>128	>128	>128	<b>Colistin</b>	4.00	4.00	4.00

### Supplemental Figure 4. MICs of PPMOs and Antibiotics used in Synergy assays.

Three strains of *P. aeruginosa*, PAO1, M57-15, and W43532 were utilized to examine if PPMOs could act favorably with current therapeutic antibiotics (Figure 5); these data are the individual MICs of compounds tested. Data shown here is representative of 3 independent experiments.