

**Table S1: PPMOs Information.** *Acyl carrier protein* (AcpP-PPMOs) are listed in the order of the location on the gene as in **Figure 1** and control (Ctrl-PPMOs) are below. The A of ATG designates 1 for the location on gene, moving 5' begins -1. Non-canonical peptides are X – 6-amino-hexanoic acid and B – beta-alanine. TEG denotes triethylene glycol.

<i>Bcc</i>						
PPMO ID	Target	PMO Sequence (5'->3')	DNA Target (5'->3')	Location on Gene	5' of PMO	3' of PMO
AcpP-0062	AcpP	TTT ACA AGT GC	GCA CTT GTA AA	-35 to -25	TEG	(RFF) <sub>3</sub> RXB
AcpP-0063	AcpP	CCT CCG AGG GA	TCC CTC GGA GG	-15 to -5	TEG	(RFF) <sub>3</sub> RXB
AcpP-0790	AcpP	CAT TAC CCC TC	GAG GGG TAA TG	-8 to 3	(RFF) <sub>3</sub> RXB	H
AcpP-0791	AcpP	CCA TTA CCC CT	AGG GGT AAT GG	-7 to 4	(RFF) <sub>3</sub> RXB	H
AcpP-0792	AcpP	TCC ATT ACC CC	GGG GTA ATG GA	-6 to 5	(RFF) <sub>3</sub> RXB	H
AcpP-0070	AcpP	GTC CAT TAC CC	GGG TAA TGG AC	-5 to 6	(RFF) <sub>3</sub> RXB	H
AcpP-0077	AcpP	GTC CAT TAC CC	GGG TAA TGG AC	-5 to 6	TEG	(RFF) <sub>3</sub> RXB
AcpP-0793	AcpP	TGT CCA TTA CC	GGT AAT GGA CA	-4 to 7	(RFF) <sub>3</sub> RXB	H
AcpP-0794	AcpP	TTG TCC ATT AC	GTA ATG GAC AA	-3 to 8	(RFF) <sub>3</sub> RXB	H
AcpP-0795	AcpP	GTT GTC CAT TA	TAA TGG ACA AC	-2 to 9	(RFF) <sub>3</sub> RXB	H
AcpP-0796	AcpP	TGT TGT CCA TT	AAT GGA CAA CA	-1 to 10	(RFF) <sub>3</sub> RXB	H
AcpP-0797	AcpP	ATG TTG TCC AT	ATG GAC AAC AT	1 to 11	(RFF) <sub>3</sub> RXB	H
AcpP-0064	AcpP	ACA CGT TGT TC	GAA CAA CGT GT	13 to 23	TEG	(RFF) <sub>3</sub> RXB
AcpP-0065	AcpP	AGT TCA GCG AC	GTC GCT GAA CT	34 to 44	TEG	(RFF) <sub>3</sub> RXB

Ctrl-0071	Control	ATC GTT GCA TC	GAT GCA ACG AT	N/A	(RFF) <sub>3</sub> RXB	H
Ctrl-0394	Control	ATC GTT GCA TC	GAT GCA ACG AT	N/A	TEG	(RFF) <sub>3</sub> RXB
Ctrl-0437	Control	ATC GTT GCA TC	GAT GCA ACG AT	N/A	(RFF) <sub>3</sub> RG	H

*P. aeruginosa*

AcpP-0445	AcpP	CTC ATA CCT TG	CAA GGT ATG AG	-6 to +5	(RXR) <sub>4</sub> XB	H
Ctrl-0078	Scramble	ATC GTT GCA TC	GAT GCA ACG AT	N/A	(RXR) <sub>4</sub> XB	H

**Table S2: *Burkholderia* Strain Information.**

<i>Burkholderia</i> Species	Isolation Source	Underlying disease
<b>B. ambifaria (AU0212)</b>		CF
<b>B. ambifaria (AU0216)</b>		CF
<b>B. ambifaria (BAA-244)</b>	Pea rhizosphere, soil	N/A
<b>B. cenocepacia (1513)</b>	Lung biopsy	CGD
<b>B. cenocepacia (3B)</b>	Lung biopsy	CGD
<b>B. cenocepacia (BAA-245)</b>		CF Patient, ET12
<b>B. cenocepacia (BC7)</b>	Sputum	CF Patient, ET12
<b>B. cenocepacia (HI2424)</b>	Soil	N/A
<b>B. cenocepacia (HI4277)</b>		CF
<b>B. cenocepacia (J2315)</b>	Sputum	CF patient, ET12
<b>B. cenocepacia (K56-2)</b>		CF patient, ET12
<b>B. cenocepacia GIIIb (AU1054)</b>	Blood	CF
<b>B. cenocepacia GIIIa (HI2718)</b>		CF Patient, ET12
<b>B. cepacia (1753)</b>	Human clinical	unknown
<b>B. cepacia (1820)</b>	Human clinical	unknown
<b>B. cepacia (1840-1)</b>	Human clinical	unknown
<b>B. cepacia (1840-2)</b>	Human clinical	unknown
<b>B. cepacia (1882)</b>	Human clinical	unknown
<b>B. cepacia (1936)</b>	Human clinical	unknown
<b>B. cepacia (2294)</b>	Human clinical	unknown
<b>B. cepacia (X49118)</b>	Blood	leukemia
<b>B. dolosa (AU0158)</b>		CF
<b>B. dolosa (AU0645)</b>		CF
<b>B. gladioli (AU15914)</b>		CF
<b>B. gladioli (HI2137)</b>	Environmental, soil	N/A
<b>B. glumae (AU6208)</b>		CF
<b>B. multivorans (AU0062)</b>		CF
<b>B. multivorans (AU0064)</b>		CF
<b>B. multivorans (BAA-247)</b>		CF
<b>B. multivorans (HI2229)</b>	Environmental, soil	N/A
<b>B. multivorans (SH-1)</b>	Lung biopsy	CGD
<b>B. multivorans (SH-2)</b>	Lung biopsy	CGD
<b>B. pyrrocinia (AU1114)</b>		CF
<b>B. pyrrocinia (AU2419)</b>		CF
<b>B. stabilis (AU26756)</b>	Human clinical	Non-CF patient
<b>B. stabilis (HI2210)</b>		CF
<b>B. vietnamiensis (AU0109)</b>		CF
<b>B. vietnamiensis (AU8819)</b>		CF
<b>B. vietnamiensis (BAA-248)</b>	Rhizosphere, soil	N/A