

## SUPPLEMENTARY INFORMATION

### ***In vitro* and *in vivo* screening for novel essential cell-envelope proteins in *Pseudomonas aeruginosa***

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**Table S1.** Plasmids used in this study.

Plasmid	Description	Reference/source
pBluescript II KS+	Cloning vector; ColE1 replicon; Ap <sup>R</sup>	Stratagene
pDM4	Suicide vector; <i>sacBR</i> , <i>oriR6K</i> ; Cm <sup>R</sup>	<sup>38</sup>
pDM4ΔPA0517	pDM4 derivative for PA0517 in-frame deletion	This study
pDM4ΔPA1645	pDM4 derivative for PA1645 in-frame deletion	This study
pDM4ΔPA1981	pDM4 derivative for PA1981 in-frame deletion	This study
pDM4ΔPA2614	pDM4 derivative for PA2614 ( <i>lolA</i> ) in-frame deletion	This study
pDM4ΔPA3786	pDM4 derivative for PA3786 in-frame deletion	This study
pDM4ΔPA4460	pDM4 derivative for PA4460 ( <i>lptH</i> ) in-frame deletion	This study
pDM4ΔPA4485	pDM4 derivative for PA4485 in-frame deletion	This study
pDM4ΔPA5126	pDM4 derivative for PA5126 in-frame deletion	This study
mini-CTX1	Self-proficient integration vector with <i>tet</i> , Ω- <i>FRT</i> - <i>attP</i> -MCS, <i>ori</i> , <i>int</i> , and <i>oriT</i> ; Tc <sup>R</sup>	<sup>43</sup>
mini-CTX1- <i>araCP<sub>BAD</sub>tolB</i>	mini-CTX1 derivative carrying <i>araCP<sub>BAD</sub>tolB</i>	<sup>19</sup>
mini-CTX1- <i>araCP<sub>BAD</sub>PA2614</i>	mini-CTX1- <i>araCP<sub>BAD</sub>tolB</i> derivative in which <i>tolB</i> has been replaced with PA2614 ( <i>lolA</i> )	This study
mini-CTX1- <i>araCP<sub>BAD</sub>PA4460</i>	mini-CTX1- <i>araCP<sub>BAD</sub>tolB</i> derivative in which <i>tolB</i> has been replaced with PA4460 ( <i>lptH</i> )	This study

Additional reference (not included in the main text):

43. Hoang, T. T., Kutchma, A. J., Becher, A. & Schweizer, H. P. Integration-proficient plasmids for *Pseudomonas aeruginosa*: site-specific integration and use for engineering of reporter and expression strains. *Plasmid* **43**, 59-72 (2000).

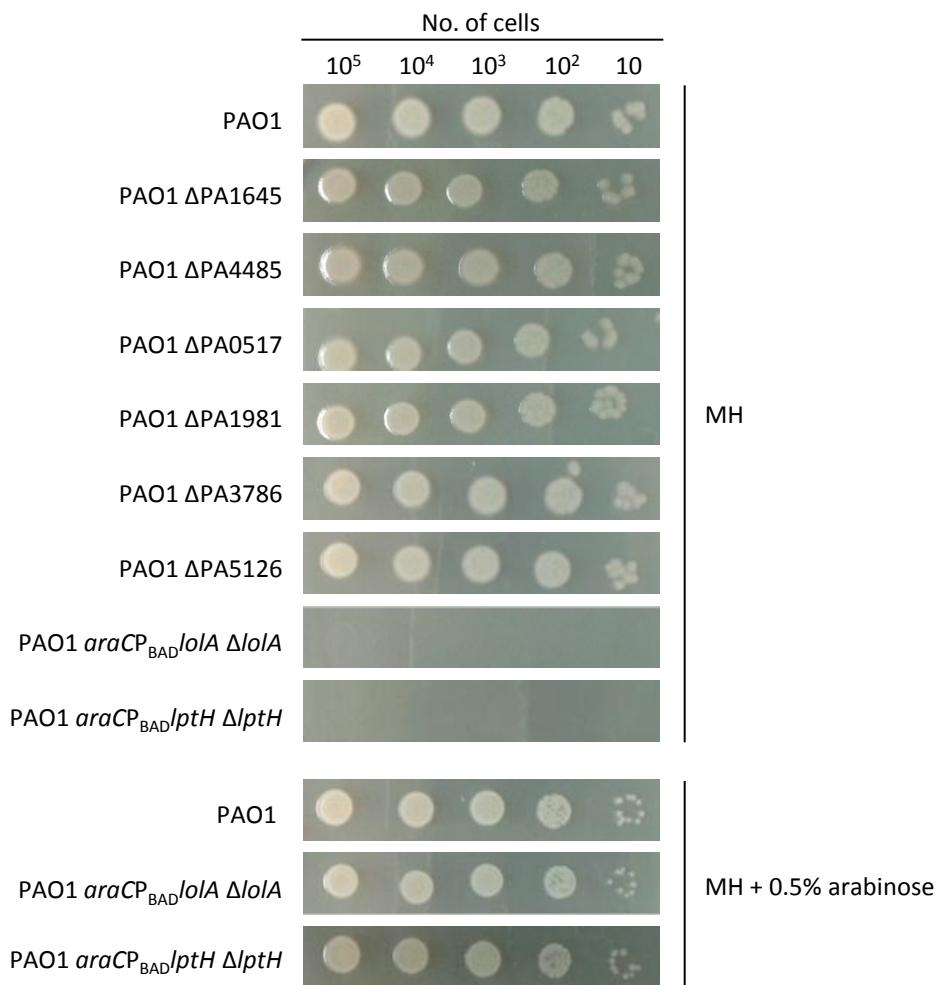
**Table S2.** Primers used in this study<sup>a</sup>

Primer name	Sequence (5'→3') <sup>b</sup>	Restriction site	Application
PA2614_FW	CCCG <u>ATATCGATGCGACTGATCCGCAC</u>	EcoRV	Generation of mini-CTX1- <i>araCP<sub>BAD</sub></i> PA2614
PA2614_RV	CGGA <u>ATTCTTACTCCTGGATCACGTCG</u>	EcoRI	Generation of mini-CTX1- <i>araCP<sub>BAD</sub></i> PA2614
PA2614mut_UP_FW	CCG <u>CTGAGGGGGCGCCGAACAAAC</u>	XhoI	Generation of pDM4ΔPA2614
PA2614mut_UP_RV	CGGG <u>ATCCCATGCCAGGGCGGC</u>	BamHI	Generation of pDM4ΔPA2614
PA2614mut_DOWN_FW	CGGG <u>ATCCTCGACGTTCCGCCGG</u>	BamHI	Generation of pDM4ΔPA2614
PA2614mut_DOWN_RV	GCT <u>CTAGACATAACCGCGAGCCCG</u>	XbaI	Generation of pDM4ΔPA2614
PA4460_FW	CCCA <u>AGCTTATGAGGTTCGTTAACCCCTC</u>	HindIII	Generation of mini-CTX1- <i>araCP<sub>BAD</sub></i> PA4460
PA4460_RV	CGGA <u>TTCTGGGCTTGAGCGTTGCC</u>	EcoRI	Generation of mini-CTX1- <i>araCP<sub>BAD</sub></i> PA4460
PA4460mut_UP_FW	ACGCG <u>TGACTACTACTGGAACGTCGGG</u>	Sall	Generation of pDM4ΔPA4460
PA4460mut_UP_RV	CGGG <u>ATCCGAGGGCGGCAGTCAG</u>	BamHI	Generation of pDM4ΔPA4460
PA4460mut_DOWN_FW	CGGG <u>ATCCCAGTAATGGCAACGCTC</u>	BamHI	Generation of pDM4ΔPA4460
PA4460mut_DOWN_RV	GCT <u>CTAGATCTGCTTGATGTCGCCG</u>	XbaI	Generation of pDM4ΔPA4460
PA0517mut_UP_FW	CGG <u>CTCGAGTACAACAAAGCGCGGC</u>	XhoI	Generation of pDM4ΔPA0517
PA0517mut_UP_RV	CGGG <u>ATCCAAGGCGTGGCTGGCGG</u>	BamHI	Generation of pDM4ΔPA0517
PA0517mut_DOWN_FW	CGGG <u>ATCCGGCTGGCTGGTGGAC</u>	BamHI	Generation of pDM4ΔPA0517
PA0517mut_DOWN_RV	GCT <u>CTAGAAGTCGGCGATCCAGATC</u>	XbaI	Generation of pDM4ΔPA0517
PA1645mut_UP_FW	CGG <u>CTCGAGCACCTGCTCCACCTCG</u>	XhoI	Generation of pDM4ΔPA1645
PA1645mut_UP_RV	CGGG <u>ATCCAGCAACAGCGGCAATACG</u>	BamHI	Generation of pDM4ΔPA1645
PA1645mut_DOWN_FW	CGGG <u>ATCCGGTCGACGAGTGCCTGC</u>	BamHI	Generation of pDM4ΔPA1645
PA1645mut_DOWN_RV	GCT <u>CTAGAGGGCGCCAGCAACCTG</u>	XbaI	Generation of pDM4ΔPA1645
PA1981mut_UP_FW	CC <u>CTCGACAAACATCAC</u>	Sall	Generation of pDM4ΔPA1981
PA1981mut_UP_RV	CGGG <u>ATCCGGCGCCGAGCAGGG</u>	BamHI	Generation of pDM4ΔPA1981
PA1981mut_DOWN_FW	CGGG <u>ATCCAAGGATTGCCCGCGGG</u>	BamHI	Generation of pDM4ΔPA1981
PA1981mut_DOWN_RV	GCT <u>CTAGAACGCTTGCAGGCGGTG</u>	XbaI	Generation of pDM4ΔPA1981
PA3786mut_UP_FW	CCG <u>CTCGAGGGTGATGCGCTGCTCG</u>	XhoI	Generation of pDM4ΔPA3786
PA3786mut_UP_RV	CGGG <u>ATCCAGCCAGGCCACGGGG</u>	BamHI	Generation of pDM4ΔPA3786
PA3786mut_DOWN_FW	CGGG <u>ATCCTGTCGCCTGACTCCTC</u>	BamHI	Generation of pDM4ΔPA3786
PA3786mut_DOWN_RV	GCT <u>CTAGATTCCACCGGCACGTCG</u>	XbaI	Generation of pDM4ΔPA3786
PA4485mut_UP_FW	ACGCG <u>TCGACTACCGTTACTTCCCC</u>	Sall	Generation of pDM4ΔPA4485

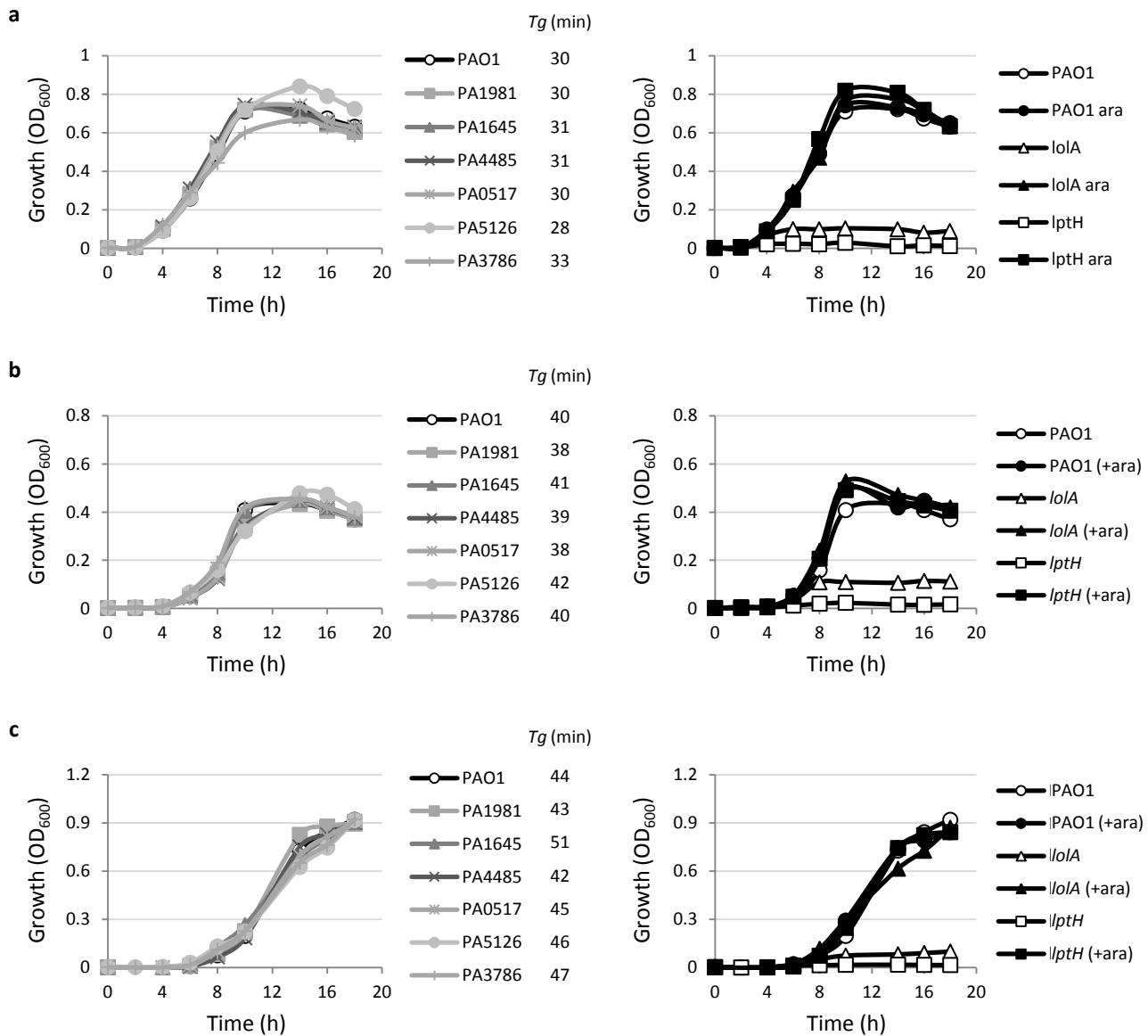
PA4485mut_UP_RV	<u>CGGGATCC</u> GGCGGATGGGGTGGC	<i>BamHI</i>	Generation of pDM4ΔPA4485
PA4485mut_DOWN_FW	<u>CGGGATCC</u> AGCGGCGTGGCGCCG	<i>BamHI</i>	Generation of pDM4ΔPA4485
PA4485mut_DOWN_RV	GCT <u>CTAGAC</u> CTGGTCACCGAAGAGGG	<i>XbaI</i>	Generation of pDM4ΔPA4485
PA5126mut_UP_FW	ACGCG <u>TCGAC</u> GCCGTGGTCCAG	<i>SalI</i>	Generation of pDM4ΔPA5126
PA5126mut_UP_RV	<u>CGGGATCC</u> GGAGGCCCTGCATGTT	<i>BamHI</i>	Generation of pDM4ΔPA5126
PA5126mut_DOWN_FW	<u>CGGGATCC</u> GACGGCCGGGAATAAAAG	<i>BamHI</i>	Generation of pDM4ΔPA5126
PA5126mut_DOWN_RV	GCT <u>CTAGAC</u> CGTCCGCAGCACAGCC	<i>XbaI</i>	Generation of pDM4ΔPA5126

<sup>a</sup> All PCRs were performed using the genomic DNA of *P. aeruginosa* PAO1 as the template.

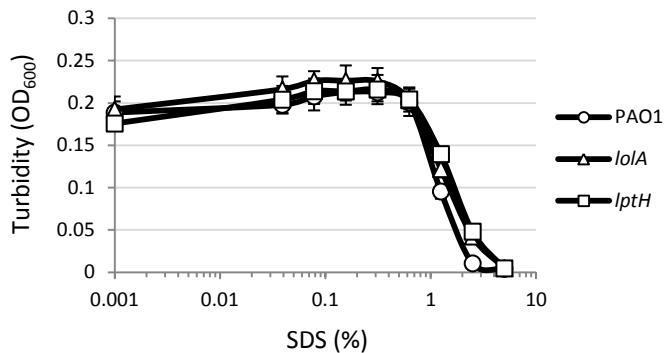
<sup>b</sup> The restriction site used for cloning is underlined in the primer sequence.



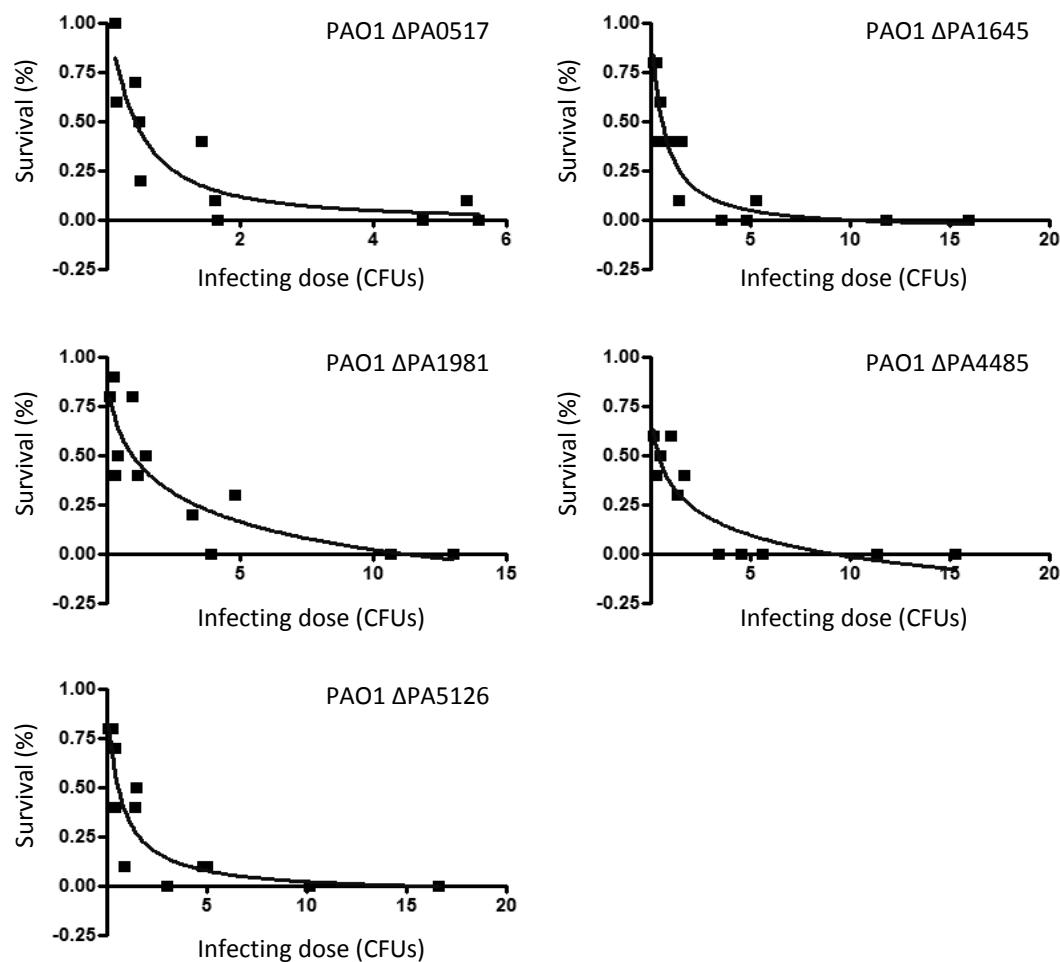
**Figure S1.** Growth of *P. aeruginosa* PAO1 and all the deletion or conditional mutants generated in this study on MH agar plates supplemented or not with 0.5% arabinose. For each strain, 5  $\mu$ l of 10-fold serial dilutions of exponential phase cultures in MH (or MH with 0.5% arabinose in the case of conditional mutants), ranging from ca.  $10^5$  to 10 viable cells, were spotted onto the plates, that were then incubated for 20 h at 37°C. The images are representative of three independent experiments giving comparable results.



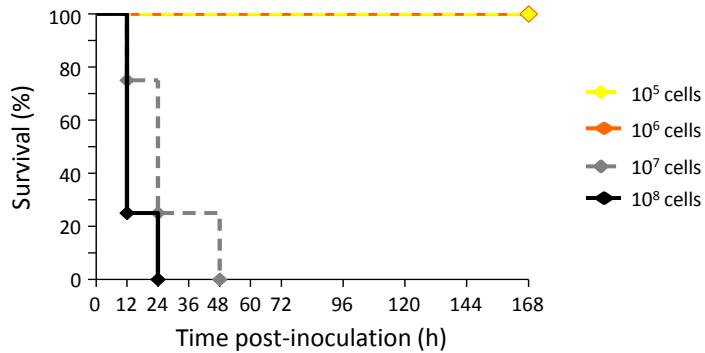
**Figure S2.** Growth curves of the wild type strain PAO1 and the PA0517, PA1645, PA1981, PA3786, PA4485 and PA5126 deletion mutants (left panels) or the *lptH* and *lolA* conditional mutants (right panels) in LB broth (**a**) or in M9 minimal medium supplemented with either succinate (**b**) or glucose (**c**) as carbon sources, at 37°C in microtiter plates at 200 rpm. Conditional mutants were cultured in the absence or in the presence of 0.5% arabinose (+ara). Generation times ( $T_g$ ) for the wild type strain and each deletion mutant are reported in the figure. Results are the mean of two independent experiments performed in triplicate, with standard deviations (SD) being < 11% of the values.



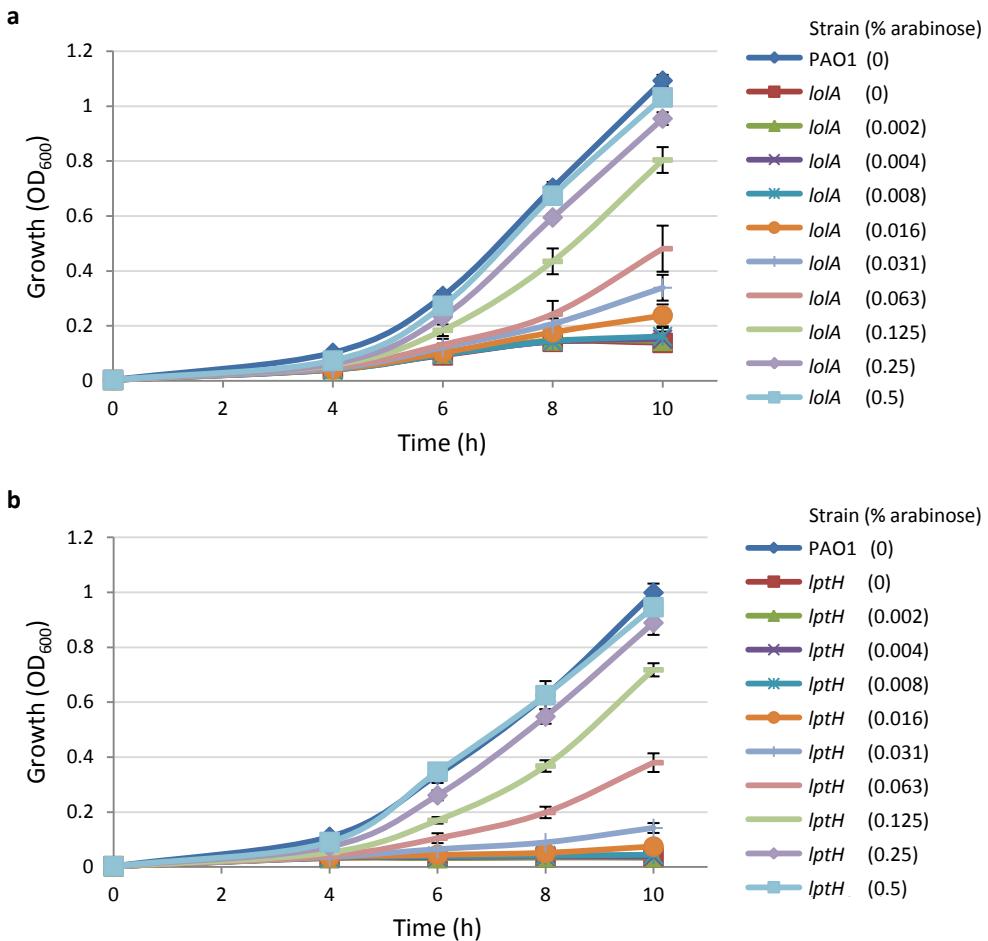
**Figure S3.** Lytic effect of different SDS concentrations (0-5%), measured as decrease in cell suspension turbidity (OD<sub>600</sub>), on the PAO1 wild type and the *lptH* and *lolA* conditional mutant grown to late-exponential phase in MH supplemented with 0.5% arabinose. Results are the mean ( $\pm$  SD) of two independent experiments performed in duplicate.



**Figure S4.** Survival curves, generated by the GraphPad Prism software, of *G. mellonella* larvae infected with different doses of the deletion mutants PAO1 ΔPA0517, PAO1 ΔPA1645, PAO1 ΔPA1981, PAO1 ΔPA4485 and PAO1 ΔPA5126.



**Figure S5.** Mortality curves for mice ( $n = 4$ ) infected with  $10^5$ ,  $10^6$ ,  $10^7$  or  $10^8$  cells of the wild type strain *P. aeruginosa* PAO1.



**Figure S6.** Growth of PAO1 and (a) the *lolA* conditional mutant or (b) the *lptH* conditional mutant in microtiter plates at 200 rpm in MH broth supplemented with increasing concentrations of arabinose (0-0.5%) after 14 h at 37°C. Results are the mean ( $\pm$  SD) of two independent experiments performed in duplicate.