

SUPPLEMENTARY INFORMATION

Mutational analysis of the essential lipopolysaccharide-transport protein LptH of *Pseudomonas aeruginosa* to uncover critical oligomerization sites

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Table S1. Frequency of spontaneous mutants resistant to ofloxacin or gentamicin in the *P. aeruginosa* *lptH* conditional mutant carrying the indicated constructs.

Construct	Frequency of spontaneous resistant mutants ^a	
	Ofloxacin	Gentamicin
pME <i>lptH</i>	3.3×10 ⁻⁷	1.9×10 ⁻⁷
pME <i>lptH</i> _β16mut	1.7×10 ⁻⁷	2.6×10 ⁻⁷
pME <i>lptH</i> _β13mut	3.0×10 ⁻⁷	1.6×10 ⁻⁷

^a Calculated from three independent assays.

Table S2. Primers used in this work.

Primer	Sequence (5'-3') ^a	Restriction sites ^b	Application
<i>lptH</i> _FW	<u>cggaattc</u> ATGCACCTGCTGTCCAACG	EcoRI	Generation of pBS
<i>lptH</i> _RV	ccgctc <u>gag</u> tcaatggatggatggatggatCTGGGCC TTTTTCTTCGGC	XhoI	<i>lptH</i>
<i>lptH</i> _β16del_FW	CAGCCGAAGAAAAAGGCCAG		Generation of β
<i>lptH</i> _β16del_RV	GTCGATGCGTGGGCGCGG		<i>lptH</i> _del1
<i>lptH</i> _β16mut_FW	acgaCAGCCGAAGAAAAAGGCCAG		Generation of pBS
<i>lptH</i> _β16mut_RV	ggtcgGTCGATGCGTGGGCGCGG		<i>lptH</i> _mut1
<i>lptH</i> _β13del_FW	GAAGGCGAGAAGATCGTCTAC		Generation of pBS
<i>lptH</i> _β13del_RV	GCCTTCCTGGATCACCTTGG		<i>lptH</i> _del2
<i>lptH</i> _β13mut_FW	agctGAAGGCGAGAAGATCGTCTAC		Generation of pBS
<i>lptH</i> _β13mut_RV	ggggcGCCTTCCTGGATCACCTTGG		<i>lptH</i> _mut2
<i>lptH</i> _β1del_FW	CAGGCCGACAGCGCCGAA		Generation of pBS
<i>lptH</i> _β1del_RV	CGGTTGTTCGCGGTCCGAAG		<i>lptH</i> _del3
<i>lptH</i> _β1mut_FW	tgatCAGGCCGACAGCGCCGAA		Generation of pBS
<i>lptH</i> _β1mut_RV	ggttcCGGTTGTTCGCGGTCCGAAG		<i>lptH</i> _mut3
<i>lptH</i> _β2del_FW	GACGACAAGCAGGGCGTCG		Generation of pBS
<i>lptH</i> _β2del_RV	GCTGTGCGCCTGGACGCG		<i>lptH</i> _del4
<i>lptH</i> _β2mut_FW	tcgcGACGACAAGCAGGGCGTCG		Generation of pBS
<i>lptH</i> _β2mut_RV	ggacgGCTGTGCGCCTGGACGCG		<i>lptH</i> _mut4
<i>lptH</i> _β15del_FW	GGTCGTGCCACCGGCTCC		Generation of pBS
<i>lptH</i> _β15del_RV	GATCTGGCGCTGGGTGTCTG		<i>lptH</i> _del5
<i>lptH</i> _β15mut_FW	tgatGGTCGTGCCACCGGCTCC		Generation of pBS
<i>lptH</i> _β15mut_RV	ggttcGATCTGGCGCTGGGTGTCTG		<i>lptH</i> _mut5
<i>lptH</i> _β16del_CHECK	CTTCGGCTGGTCGATGCG		Colony PCR
<i>lptH</i> _β16mut_CHECK	CGGCTGTCGTGGTCGGTC		screening of pBS
<i>lptH</i> _β13del_CHECK	TCGCCTTCGCCTTCCTGG		constructs with the
<i>lptH</i> _β13mut_CHECK	CTCGCCTTCAGCTGGGGC		different <i>lptH</i>
<i>lptH</i> _β1del_CHECK	GTCGGCCTGCGGTTGTTC		variants (each
<i>lptH</i> _β1mut_CHECK	TGTCGGCCTGATCAGGTTT		reverse primer was
<i>lptH</i> _β2del_CHECK	GCTTGTCGTGCTGTCTGG		coupled with
<i>lptH</i> _β15del_CHECK	TGGCACGACCGATCTGGC		<i>lptH</i> _FW)
<i>lptH</i> _β15mut_CHECK	TGGCACGACCATCAGGTTT		
<i>lptH</i> _pET28b_FW	ggaattc <u>cata</u> TGCCTTCGGACCGCGAAC	NdeI	Generation of
<i>lptH</i> _pET28b_RV	cccaagctt <u>tca</u> CTGGGCCTTTTCTTCGGC	HindIII	pET28b <i>lptH</i> and mutant constructs
<i>recA</i> _mut_UP_FW	ccgcTCGAGGCTGGCTACGTGAC	XhoI	Generation of
<i>recA</i> _mut_UP_RV	cgggatCCGAATTGGCGTTTCGATCTG	BamHI	pDM4 Δ <i>recA</i>
<i>recA</i> _mut_DOWN_FW	cgggatCCGAAGAAGTGGCTGACGC	BamHI	
<i>recA</i> _mut_DOWN_RV	gctctaGAAAACTGGGACTGAAGCTG	XbaI	
M13_FW	GTTTTCCAGTCACGAC		DNA sequencing
M13_RV	AACAGCTATGACCATG		
T7_FW	TAATACGACTCACTATAGGG		

^a Lowercase letters indicate the region of the primer that does not anneal to the template.^b The restriction site used for cloning is underlined in the primer sequence.

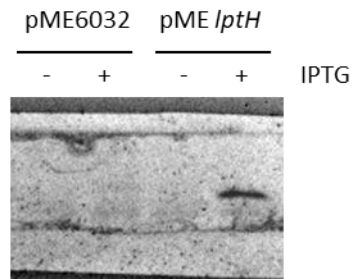


Figure S1. Detection of LptH in the *P. aeruginosa* *lptH* conditional mutant carrying the empty vector pME6032 or pME *lptH* cultured in MH containing 0.5% arabinose and supplemented (+) or not (-) with 0.5 mM IPTG, by Western blotting of whole-cell lysates (20 µg total proteins) with an anti-LptH polyclonal antibody. The image is representative of two independent experiments which gave similar results.

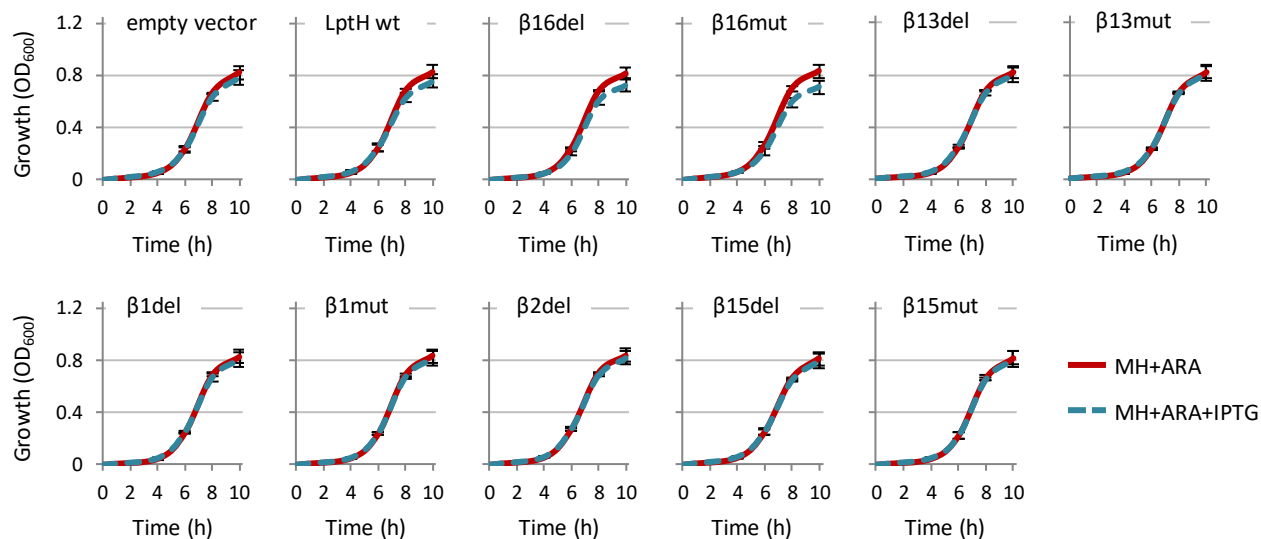


Figure S2. Growth curves of the *P. aeruginosa* *lptH* conditional mutant carrying the empty plasmid pME6032, pME6032 with wild-type *lptH* or pME6032 with different mutant variants in MH at 37°C in microtiter plates in the presence of 0.5% arabinose (+ ARA; red lines) or 0.5% arabinose and 0.5 mM IPTG (+ ARA + IPTG; dashed blue lines). Growth was measured as OD₆₀₀. Results are the mean ($\pm$ SD) of two independent experiments, each performed in triplicate.

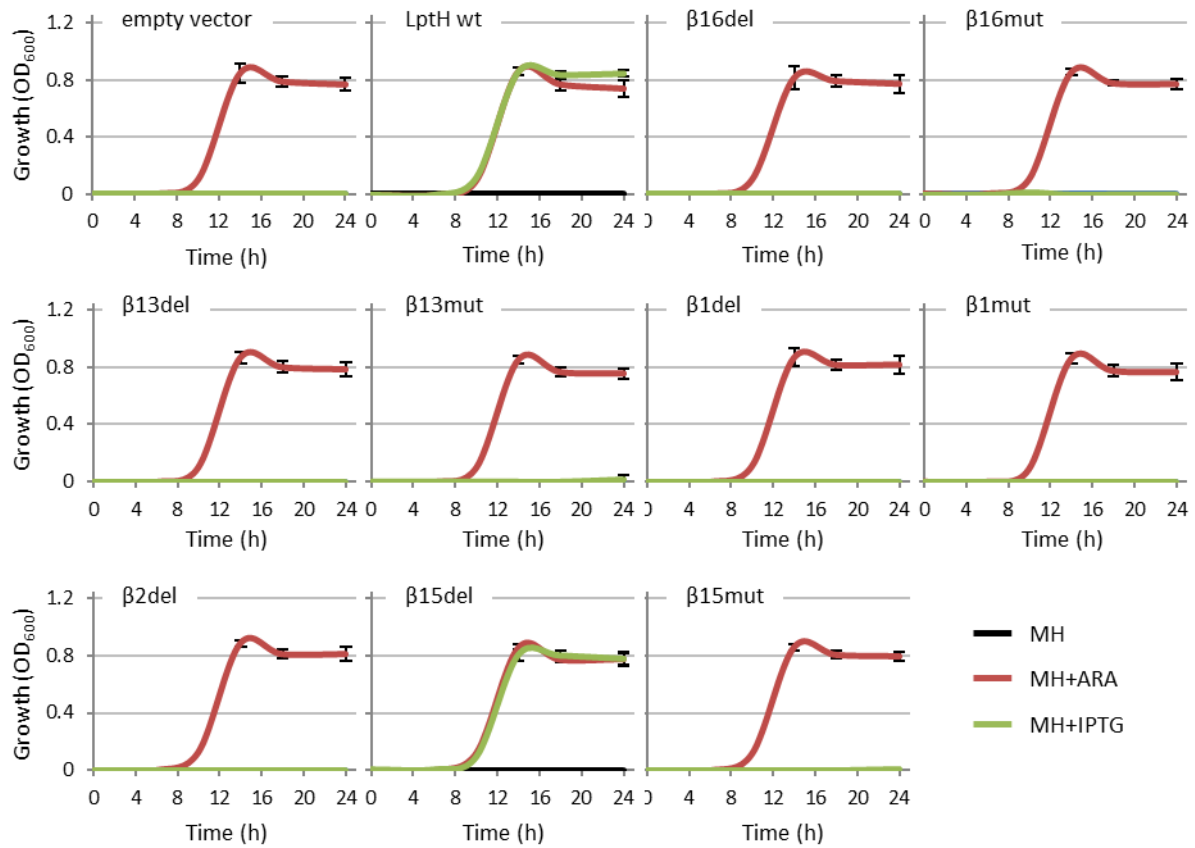


Figure S3. Growth curves of the *P. aeruginosa* *lptH* conditional mutant carrying the empty plasmid pME6032, pME6032 with wild-type *lptH* or pME6032 with different mutant variants inoculated at a cell density of ca. 50-100 cells/mL in MH at 37°C in microtiter plates in the absence (black lines) or in the presence of 0.5% arabinose (+ ARA; red lines) or 0.5 mM IPTG (+ IPTG; green lines). Growth was measured as OD₆₀₀. Results are the mean ($\pm$ SD) of three independent experiments, each performed in quadruplicate.

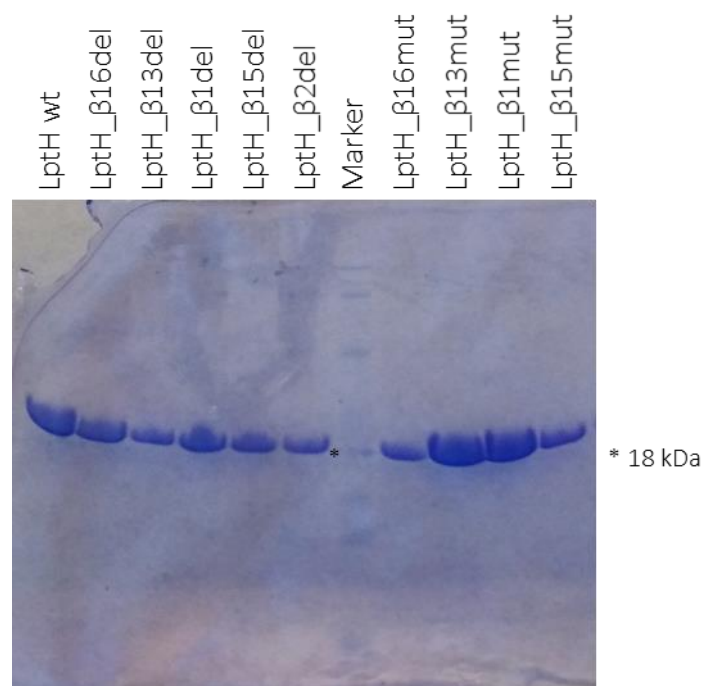


Figure S4. Coomassie-stained SDS-PAGE gel showing the proteins purified in this work.

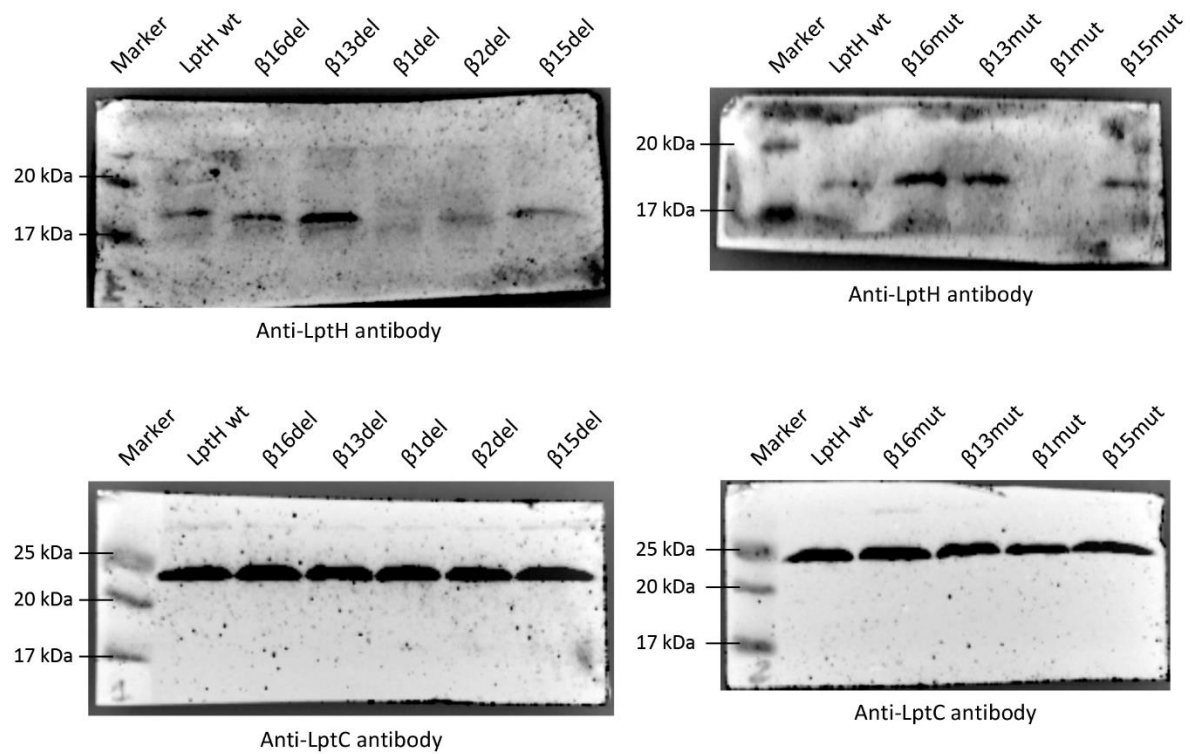


Figure S5. Full-length blots for the cropped images shown in Figure 2B.