

The *Pseudomonas aeruginosa* DksA1 protein is involved in H₂O₂ tolerance and within-macrophages survival and can be replaced by DksA2

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SUPPLEMENTAL MATERIAL

Figure S1. Effect of *dksA1 dksA2* double deletion on phagocytosis

Figure S2. Expression analysis of *dksA1* and *dksA2* in response to H₂O₂

Figure S3. Expression analysis of *katN*

Table S1. Bacterial strains used in this study

Table S2. Oligonucleotides used in this study

Figure S1

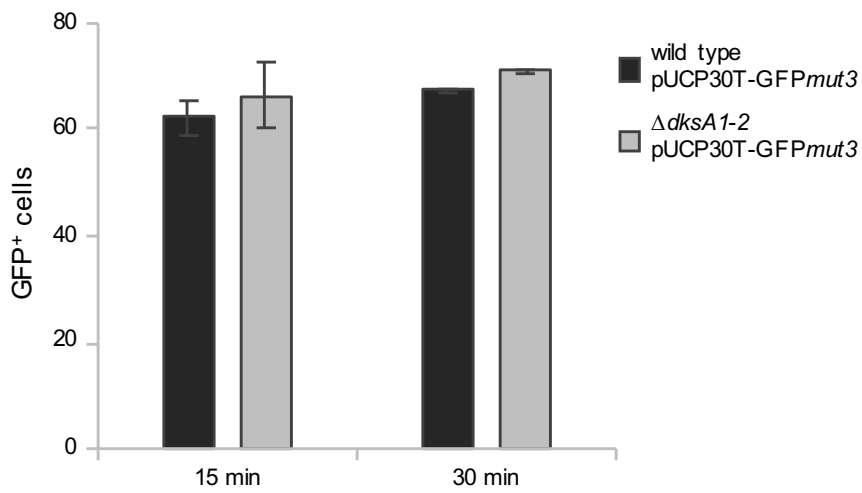


Figure S1. Effect of *dksA1 dksA2* double deletion on phagocytosis. Fluorescent macrophages in cell populations infected with the indicated GFP-expressing strains for 15 or 30 min. The average of two independent experiments, each in duplicate, is shown with standard deviations.

Figure S2

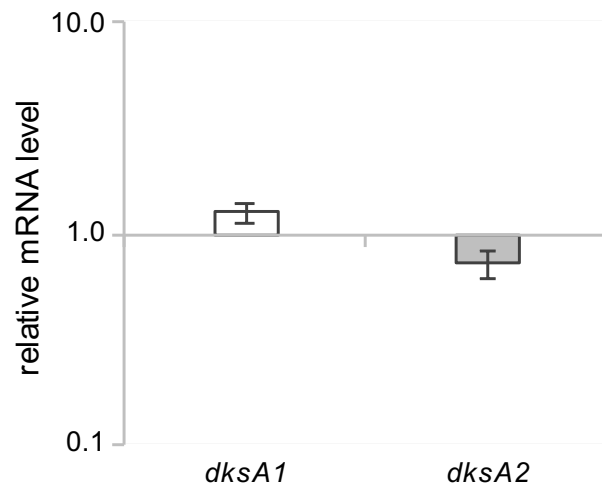


Figure S2. Expression analysis of *dksA1* and *dksA2* in response to H₂O₂. Real Time PCR analysis showing the relative mRNA level of the *dksA1* and *dksA2* genes. *P. aeruginosa* PAO1 grown to the exponential phase (OD₆₀₀ ≈ 2.5) in LB with 0.1 mM IPTG was challenged with 1 mM H₂O₂ for 20 min. Gene expression is reported as fold change relative to the untreated culture. The average of two independent analyses, each performed on three technical replicates, is shown with standard deviations.

Figure S3

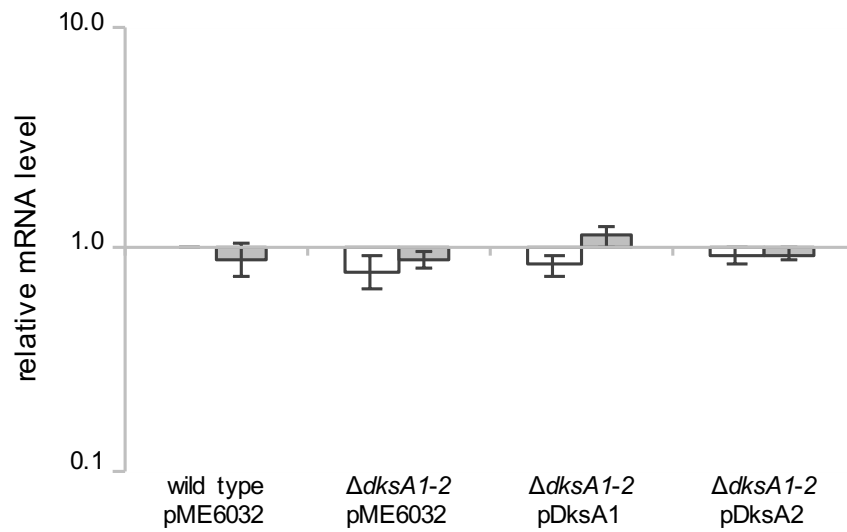


Figure S3. Expression analysis of *katN*. Bacterial cultures were grown to the exponential phase ($OD_{600} \approx 2.5$) in LB supplemented with 0.1 mM IPTG and then incubated with or without 1 mM H_2O_2 for 20 min. The mRNA levels of *katN* gene were determined by Real Time PCR analysis. Gene expression in the indicated untreated (white bars) or treated (grey bars) strains is reported as fold-change in gene expression relative to the untreated wild type PAO1(pME6032) strain. The average of two independent analyses, each performed on three technical replicates, is shown with standard deviations.

Table S1. Bacterial strains used in this study.

Strain	Relevant characteristics	Reference/Source
<i>Pseudomonas aeruginosa</i>		
PAO1(pME6032)	Wild type strain (ATCC 15692) containing the pME6032 expression vector.	[24]
PAO1(pUCP30T-GFP <i>mut3</i>)	Wild type strain (ATCC 15692) containing the pUCP30T-GFP <i>mut3</i> vector for the GFP constitutive expression [50].	This study
PAO1 $\Delta dksA1-2$ (pME6032)	PAO1 derivative strain carrying a deletion of the <i>dksA1</i> and <i>dksA2</i> genes and containing the pME6032 expression vector.	[24]
PAO1 $\Delta dksA1-2$ (pUCP30T-GFP <i>mut3</i>)	PAO1 derivative strain carrying a deletion of the <i>dksA1</i> and <i>dksA2</i> genes and containing the pUCP30T-GFP <i>mut3</i> vector for the GFP constitutive expression [50].	This study
PAO1 $\Delta dksA1-2$ (pDksA1)	PAO1 derivative strain carrying a deletion of the <i>dksA1</i> and <i>dksA2</i> genes and containing a pME6032 derivative for the IPTG-dependent expression of <i>dksA1</i> .	[24]
PAO1 $\Delta dksA1-2$ (pDksA2)	PAO1 derivative strain carrying a deletion of the <i>dksA1</i> and <i>dksA2</i> genes and containing a pME6032 derivative for the IPTG-dependent expression of <i>dksA2</i> .	[24]

Table S2. Oligonucleotides used in this study.

Name	Sequence (5'→3')^a	Restriction site^b
FW <i>Ppq</i> sL	TCCGCTCGAGGATCGTCACCGTCAACTG	XhoI
RVP <i>Ppq</i> sL	TAACTGCAGCGTCATGGATGAGTCTCCG	PstI
FW <i>dksA</i> 1RT	TTCACCGCGATCCTGAACAA	-
RV <i>dksA</i> 1RT	ATCAGTTGCAGGGTCTCGTC	-
FW <i>dksA</i> 2RT	AAGCCCAGCAGGACTTCTTC	-
RV <i>dksA</i> 2RT	TGTCGAGCAGCTTCTTCTCC	-
FW <i>kat</i> ART	ATGCGTTTCTACACCGAGCA	-
RV <i>kat</i> ART	ATGGTCAACTGATGCAGCGA	-
FW <i>kat</i> BRT	TGTATTCCAACGAGGGCACC	-
RV <i>kat</i> BRT	GCTCACCAGGTCATTGGTCA	-
FW <i>kat</i> ERT	GAGAACCAGCAGGACAGCTT	-
RV <i>kat</i> ERT	GGGTCGTTGGTGAAGTCGAT	-
FW <i>kat</i> NRT	ACTCGCATATCACCTCGCTG	-
RV <i>kat</i> NRT	AAGGTCAGCGCATCCTTGAT	-
FW <i>rpo</i> DRT	CCACCGCCGAACTCACC	-
RV <i>rpo</i> DRT	CTCGTCGGTCTCGTGGTT	-

^a The restriction sites included in the oligos are underlined.

^b -, no restriction site introduced.