

**Supplementary Table S1. Bacterial strains and plasmids used in this study**

Strains or plasmids	Relevant characteristics <sup>a</sup>	Source
<b>Strains</b>		
<i>Xanthomonas. oryzae</i> pv. <i>oryzae</i>		
PXO99 <sup>A</sup>	The <i>Xoo</i> wild-type strain, Philippine race 6	[32]
8-24	<i>minC</i> Tn5 insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This study
24-46	<i>minD</i> Tn5 insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This study
PΔ <i>minC</i>	<i>minC</i> deletion mutant of PXO99 <sup>A</sup>	This study
PΔ <i>minD</i>	<i>minD</i> deletion mutant of PXO99 <sup>A</sup>	This study
PΔ <i>minCDE</i>	Triple <i>minC</i> , <i>minD</i> and <i>minE</i> deletion mutant of PXO99 <sup>A</sup>	This study
C8-24	8-24 harbouring pML123- <i>minC</i> ; Km <sup>r</sup> , Gm <sup>r</sup>	This study
CPΔ <i>minC</i>	PΔ <i>minC</i> harbouring pML123- <i>minC</i> ; Gm <sup>r</sup>	This study
CPΔ <i>minD</i>	PΔ <i>minD</i> harbouring pML123- <i>minD</i> ; Gm <sup>r</sup>	This study
CPΔ <i>minCDE</i>	PΔ <i>minCDE</i> harbouring pML123- <i>minCDE</i> ; Gm <sup>r</sup>	This study
PΔ <i>hrpG</i>	<i>hrpG</i> deletion mutant of PXO99 <sup>A</sup>	This laboratory
PΔ <i>hrcU</i>	<i>hrcU</i> deletion mutant of PXO99 <sup>A</sup>	This laboratory
PΔ <i>trh</i>	<i>trh</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
PΔ <i>xrvA</i>	<i>xrvA</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
PΔ <i>zur</i>	<i>zur</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
PΔ <i>rpfC</i>	<i>rpfC</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
PΔ <i>rpfG</i>	<i>rpfG</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
PΔ <i>rpfF</i>	<i>rpfF</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
PΔ <i>clp</i>	<i>clp</i> insertion mutant of PXO99 <sup>A</sup> ; Km <sup>r</sup>	This laboratory
<b>Escherichia coli</b>		
DH5a	φ901ac ZΔM15, recA1	Invitrogen
<b>Plasmids</b>		
pHG2- <i>hrpF</i>	The <i>Xoo</i> <i>hrpX</i> promoter cloned in pHG2; Sp <sup>r</sup>	[33]
pHG3- <i>hrpB1</i>	The <i>Xoo</i> <i>hrpB1</i> promoter cloned in pHG3; Sp <sup>r</sup>	[36]
pHG2- <i>hrpG</i>	The <i>Xoo</i> <i>hrpG</i> promoter cloned in pHG2; Sp <sup>r</sup>	[36]
pHG2- <i>hrpX</i>	The <i>Xoo</i> <i>hrpX</i> promoter cloned in pHG2; Sp <sup>r</sup>	[36]
pH1- <i>hrpG</i> ::FLAG	The fusion of <i>hrpG</i> ::FLAG cloned in pH1; Sp <sup>r</sup>	[33]
pH3- <i>hrpX</i> ::FLAG	The fusion of <i>hrpX</i> ::FLAG cloned in pH3; Sp <sup>r</sup>	[33]
pH3- <i>hrpB1</i> ::FLAG	The fusion of <i>hrpB1</i> ::FLAG cloned in pH3; Sp <sup>r</sup>	[33]
pML123- <i>minC</i>	The <i>minC</i> complementary fragment cloned in pML123; Gm <sup>r</sup>	This study
pML123- <i>minD</i>	The <i>minD</i> complementary fragment cloned in pML123; Gm <sup>r</sup>	This study
pML123- <i>minCDE</i>	The <i>minCDE</i> complementary fragment cloned in pML123; Gm <sup>r</sup>	This study
pHM1- <i>gfp</i>	A high-copy vector harbouring a <i>gfp</i> gene derived from pHM1; Sp <sup>r</sup> , Ap <sup>r</sup>	This laboratory
pHG2- <i>hrpG</i> - <i>post</i>	The post transcriptional fusion of <i>hrpG</i> :: <i>uidA</i> cloned in pH2; Sp <sup>r</sup>	This laboratory

<sup>a</sup>Ap<sup>r</sup>, ampicillin resistance; Km<sup>r</sup>, kanamycin resistance; Gm<sup>r</sup>, gentamycin resistance; Sp<sup>r</sup>,

streptomycin resistance

**Supplementary Table S2:** Primer sequences used in this study

Primers Name	Primer Sequences (5'-3')	Description
<i>minC</i> up-F	GTGTCTAGACGCATGTAATAATCCAACGC	
<i>minC</i> up-R	ATGAAGCTTAGCACACCGGCTTCGTGCAG	For <i>minC</i> deletion mutant
<i>minC</i> down-F	GAGAAGCTTATCTTCTGCCGCGACTTCCA	construction
<i>minC</i> down-R	ACTCCCGGGCAGCCAGGTCCTTGAGCAC	
<i>minD</i> up-F	ACTCTGCAGCGATGCGGCCTTGCTGTTC	
<i>minD</i> up-R	GACTCTAGAGGCTTGCGCTGGTGGTGGT	For <i>minD</i> deletion mutant
<i>minD</i> down-F	AAGTCTAGAGCGTGGAAGGCGGCGAGATG	construction
<i>minD</i> down-R	AGAGGATCCCGGTCGGGTCACAGGCAACA	
<i>minCDE</i> up-F	AACTGCGATTATATGCCGATGTGCAGA	
<i>minCDE</i> up-R	GACTCTAGAGGTCACGCGCTCACGCATT	For <i>minC</i> , <i>minD</i> and <i>minE</i> triple
<i>minCDE</i> down-F	AAGTCTAGAGAAGACGGCGACAAGTAAG	deletion mutant construction
<i>minCDE</i> down-R	AGAGGATCCAGTGCCTCGGTTCGAACAGT	
GNAT- <i>minC</i> -F	CGGGTTTCTGGTCGGATTCG	For confirmation of GNAT- <i>minC</i>
GNAT- <i>minC</i> -R	GGAAGTCGCGGCAGAAGATG	operon
<i>minC</i> -D-F	ACCACCAGCGCAAGCCTGGC	For confirmation of <i>minC</i> -D
<i>minC</i> -D-R	GTTCGCCATCCTTGACCAGA	operon
<i>minD</i> -E-F	ACCACCAGCGCAAGCCTGGC	For confirmation of <i>minD</i> -E
<i>minD</i> -E-R	GTTCGCCATCCTTGACCAGATC	operon
<i>minE</i> - <i>PXO</i> _04466-F	ATGGGCCTGCTCGATTTTCT	For confirmation of <i>minE</i> -
<i>minE</i> - <i>PXO</i> _04466-R	TCACGCAGGCAACAAGCCGC	<i>PXO</i> _04466 operon
<i>minC</i> com-F	GAGAAGCTTGAACAACAATGCCGGACTG	For <i>minC</i> complementray fragment
<i>minC</i> com-R	GATGTCGACACCACGCGCCGCTCGCAGC	
<i>minCDE</i> com-F	GAGAAGCTTGAACAACAATGCCGGACTG	For <i>minC</i> , <i>minD</i> and <i>minE</i>
<i>minCDE</i> com-R	GATGTCGACTCGCTCGCGGTCGGGTCACAG	complementray fragment
<i>minD</i> com-F	GAGAAGCTTATCGATGTTGACGTACTTC	For <i>minD</i> complementray fragment
<i>minD</i> com-R	GATGTCGACAGTTGTATGCGGAGAACTG	
<i>hrpB1</i> -F	TTCGATGCATGGATTTTCGATCAAGC	<i>hrpB1</i> gene fragment for qRT-PCR
<i>hrpB1</i> -R	CGCCGGTGCGGACGTTGGGGTAGTT	
<i>rpfC</i> -F	ACCACGCGTCTGGATGTAGAGCAGAAGG	<i>rpfC</i> gene fragment for qRT-PCR
<i>rpfC</i> -R	GCACCTCTTCCACCAGCGACAGCAGACT	
<i>rpfG</i> -F	TGGAGCAGCGCTTGCTGGCCAGCATGAA	<i>rpfG</i> gene fragment for qRT-PCR
<i>rpfG</i> -R	CGACAACCCAGTTGCTCGGCAATCAGG	
<i>rpfF</i> -F	CAGTTGCTTGGCATGGGTCT	<i>rpfF</i> gene fragment for qRT-PCR
<i>rpfF</i> -R	TGGTACTGCGGTGGTCATTT	
<i>clp</i> -F	GTTGTTCATCGAATCCGATACCCGCGAG	<i>clp</i> gene fragment for qRT-PCR
<i>clp</i> -R	ATCTTCGGCGCATCCGGCGACAGGCTGG	
<i>xrvA</i> -F	CGAACTGCAGAAGCTGGAAGAACAGGAG	<i>xrvA</i> gene fragment for qRT-PCR
<i>xrvA</i> -R	CTTCGCTGCGGCCGCTCTTGACCACGTT	
<i>trh</i> -F	CAACTTGAAGCCGAAGGCCGCATCTACC	<i>trh</i> gene fragment for qRT-PCR

<i>trh</i> -R	ACGTACTGCATGAAACCGGCGATGCTGG	
<i>rpoD</i> qRT-F	CGACAACACCACCAACATCAATC	<i>rpoD</i> gene fragment for qRT-PCR
<i>rpoD</i> qRT-R	GCTTACCGACCTCTTCCAACG	
<i>gyrB</i> qRT-F	CGGCACTTACGACTCCAGCAAG	<i>gyrB</i> gene fragment for qRT-PCR
<i>gyrB</i> qRT-R	CGACCAGGATTTTCACCACGATG	

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Note: Underlined bases indicate restriction enzyme sites