

The *Yersinia pestis* HmsCDE regulatory system is essential for blockage of the oriental rat flea (*Xenopsylla cheopis*), a classic plague vector

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SUPPORTING INFORMATION

SUPPORTING TABLES

Table S1. Bacterial strains and plasmids used in this study

Strain or plasmid	Description	Reference or source
Strains		
<i>Escherichia coli</i>		
DH5 α	Cloning strain	Invitrogen
DH5 α (λ <i>pir</i>)	Strain for maintenance of R6K origin suicide vectors	S. C. Straley
<i>Yersinia pestis</i> ^a		
KIM5 (pCD1Ap)+	Pgm ⁺ (Hms ⁺) Lcr ⁺ (pCD1Ap; <i>yadA::amp</i>); pMT1, pPCP1; Ap ^r	(Gong et al., 2001)
KIM5-2173.1 (pCD1Ap)+	Pgm ⁺ (Hms ⁺) Δ <i>hmsC2173.1</i> Lcr ⁺ (pCD1Ap; <i>yadA::amp</i>); pMT1, pPCP1; Ap ^r	This study
KIM6+	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT1, pPCP1	(Fetherston et al., 1992)
KIM6-2051+	Pgm ⁺ Hms ⁻ (<i>hmsT2051::mini-kan</i>) Lcr ⁻ ; pMT1, pPCP1; Km ^r	(Jones et al., 1999)
KIM6-2090.2+	Pgm ⁺ Δ <i>hmsP2090.2</i> Lcr ⁻ ; pMT1, pPCP1	(Bobrov et al., 2011)
KIM6-2093+	Pgm ⁺ (Hms ⁺) Δ <i>y2360::cam2093</i> Lcr ⁻ ; pMT1, pPCP1; Cm ^r	(Bobrov et al., 2011)
KIM6-2118	Pgm ⁺ Hms ⁻ (Δ <i>hmsR2118</i>) Lcr ⁻ ; pMT1,	(Forman et al., 2006)

	pPCP1	
KIM6-2159.1+	Pgm ⁺ <i>ΔhmsD2159.1</i> Lcr ⁻ ; pMT1, pPCP1	(Bobrov et al., 2011)
KIM6-2173.1+	Pgm ⁺ <i>ΔhmsC2173.1</i> Lcr ⁻ ; pMT1, pPCP1	This study
KIM6-2173.2	Pgm ⁺ Hms ⁻ (<i>ΔhmsR2118</i>) <i>ΔhmsN2173.1</i> Lcr ⁻ ; pMT1, pPCP1	This study
KIM6-2173.3+	Pgm ⁺ <i>ΔhmsC2173.1 ΔhmsT</i> Lcr ⁻ ; pMT1, pPCP1	This study
KIM6-2173.4 +	Pgm ⁺ <i>ΔhmsC2173.1 ΔhmsT</i> <i>ΔhmsE::kan2174</i> Lcr ⁻ ; pMT1, pPCP1; Km ^r	This study
KIM6-2174+	Pgm ⁺ <i>ΔhmsE::kan2174</i> Lcr ⁻ ; pMT1, pPCP1; Km ^r	This study
KIM6-2174.1+	Pgm ⁺ <i>ΔhmsE 2174.1</i> Lcr ⁻ ; pMT1, pPCP1	This study
KIM6-2198+	Pgm ⁺ <i>ΔhmsCD2198</i> Lcr ⁻ ; pMT1, pPCP1	This study
KIM10+	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT1	(Perry et al., 1990)
CO92 ^c	Pgm ⁺ Lcr ⁻ ; pMT, pPCP	S.W. Bearden, CDC
CO92 <i>ΔhmsC</i>	Pgm ⁺ <i>ΔhmsC2173.1</i> Lcr ⁻ ; pMT, pPCP	This study
Kimberley	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT, pPCP	(Fetherston et al., 1992)
Kuma	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT, pPCP	(Perry et al., 1990)
Kuma <i>ΔhmsC</i>	Pgm ⁺ <i>ΔhmsC2173.1</i> Lcr ⁻ ; pMT, pPCP	This study
<i>Pestoides D</i>	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT	S.W. Bearden, CDC
<i>Pestoides D</i> <i>ΔhmsC</i>	Pgm ⁺ <i>ΔhmsC2173.1</i> Lcr ⁻ ; pMT	This study
<i>Pestoides F</i>	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT	S.W. Bearden, CDC

<i>Pestoides</i> F	Pgm ⁺ Δ <i>hmsC</i> 2173.1 Lcr ⁻ ; pMT	This study
Δ <i>hmsC</i>		
Salazar	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT, pPCP	(Perry et al., 1990; Fetherston et al., 1992)
Yokohama	Pgm ⁺ (Hms ⁺) Lcr ⁻ ; pMT, pPCP	(Perry et al., 1990; Fetherston et al., 1992)
Plasmids		
pBAD30	Low copy number expression vector; <i>araC</i> ⁺ ; <i>araBAD</i> promoter; Ap ^r	(Guzman et al., 1995)
pCP20	FLP recombinase expressing plasmid; Ap ^r Cm ^r	(Cherepanov and Wackernagel, 1995)
pKD3	Template plasmid; Cm ^r	(Datsenko and Wanner, 2000)
pKD13	Template plasmid; Km ^r	(Datsenko and Wanner, 2000)
pKD46	Red recombinase expressing plasmid; Ap ^r	(Datsenko and Wanner, 2000)
pRMCD28-T5	<i>phoA</i> fusion vector with T5 promoter; Ap ^r	(Bobrov et al., 2008)
pRMCD70-T5	<i>lacZ</i> fusion vector with T5 promoter; Ap ^r	(Bobrov et al., 2008)
pSkippy	FLP recombinase expressing plasmid; Ap ^r	(Price et al., 2012)
pWL204	Derivative of pKD46; Ap ^r Sac ^s	(Lathem et al., 2007)
pBAD- <i>hmsC</i>	658 bp fragment encoding <i>hmsC</i> cloned into pBAD30 by SmaI/XbaI; arabinose	This study

	inducible expression; Ap ^r	
pBAD- <i>hmsE</i>	619 bp fragment encoding <i>hmsE</i> cloned into pBAD30 by EcoRI/SmaI; arabinose inducible expression; Ap ^r	This study
pKNG- Δ <i>hmsR</i>	Fragment with Δ <i>hmsR</i> cloned into pKNG101; Sm ^r Sac ^s	(Forman et al., 2006)
pKNG- Δ <i>hmsT</i> - Δ <i>scar</i>	Fragment with Δ <i>hmsT</i> - Δ <i>scar</i> cloned into pKNG101; Sm ^r Sac ^s	(Bobrov et al., 2011)
pRMCD28-T5- <i>hmsC</i> -R207	<i>hmsC</i> -R207- <i>phoA</i> gene fusion cloned into pRMCD28-T5 by XbaI/PstI; Ap ^r	This study
pRMCD70-T5- <i>hmsC</i> -R207	<i>hmsC</i> -207- <i>lacZ</i> gene fusion cloned into pRMCD70-T5 by XbaI/PstI; Ap ^r	This study
pRMCD28-T5- <i>hmsD</i> -G425	<i>hmsD</i> -G425- <i>phoA</i> gene fusion cloned into pRMCD28-T5 by XbaI/PstI; Ap ^r	This study
pRMCD70-T5- <i>hmsD</i> -G425	<i>hmsD</i> -G425- <i>lacZ</i> gene fusion cloned into pRMCD70-T5 by XbaI/PstI; Ap ^r	This study
pRMCD28-T5- <i>hmsD</i> -R156	<i>hmsD</i> -R156- <i>phoA</i> gene fusion cloned into pRMCD28-T5 by XbaI/PstI; Ap ^r	This study
pRMCD70-T5- <i>hmsD</i> -R156	<i>hmsD</i> -R156- <i>lacZ</i> gene fusion cloned into pRMCD70-T5 by XbaI/PstI; Ap ^r	This study

^a All *Y. pestis* strains lacking the pCD plasmid (Lcr⁻) are avirulent. KIM5 strains were constructed by electroporation of pCD1Ap into KIM derivatives. KIM

derivatives with plus sign in possess an intact 102-kb chromosomal *pgm* locus which contains, among others, the *hmsHF*RS operon that is essential for biofilm formation. All KIM derivatives lacking a plus sign have either a *pgm* deletion or a mutation within the *pgm* locus.

- ^b Abbreviations: Ap^r, Cm^r, Km^r, and Sm^r, respectively indicate resistance to ampicillin, chloramphenicol, kanamycin, and streptomycin.
- ^c CO-99-3015 is the CDC stock of CO92. For simplicity, CO92 is used in the text.

Table S2. Primers used in this study.

<i>Primer name</i>	<i>Primer sequence (5'→3')</i>
<i>Gene deletion</i>	
Y3729 RED-1	CAACCACCACATTTAGCATCCCTTTTTAGCCCAACAGG AAGCGTAAAAGTGTAGGCTGGAGCTGCTTCG
Y3729 RED-2	TTCGGCTTCATCGTGCATTAGCCTTCTTACGGGCAAGT TGCAATACACATTCCGGGGATCCGTGCGACC
Y3729 RED-2-pKD3	TCGGCTTCATCGTGCATTAGCCTTCTTACGGGCAAGTT GCAATACACTACATATGAATATCCTCCTTAGT
Y3731 RED-1	TGCCTATCAGCCGAATACGACTAACACAGAAAGTTT GATAGGTGACGTGTAGGCTGGAGCTGCTTCG
Y3731 RED-2	ACAGCCGCAAACCGGTAAATAAAGGACGTTAGGACGC GGTGATAATAAATTCCGGGGATCCGTGCGACC
Y3730 RED-2	TTTGCATTTTCTGCATGTCACCTATCCTAAACTTTCTGT CATATGAATATCCTCCTTA
<i>Control of deletion</i>	
Y3729-1	TAGCCCAACAGGAAGCGTAA
Y3729-Xba-2	GATTCTAGATCGGCTTCATCGTGCATTAG
Y3730delProbeR	CACATCACTGTCTGAATAGCACT
Cm-Alex1	AATATCCAGCTGAACGGTCTG

HmsT-AO-1	GTGGTACAACATGCTGACGG
HmsT-AO-2	CACCAGAAAGGGGAAATGGG
HmsRO-XbaI	GCTCTAGACTCATGATTTACCCTCCCAAT
HmsRO-EcoRI	GGAATTCTACTGGCGAACCACCGCTAAAAG
pQE3730R	CTAAGATCTTCCTAAACTTTCTGTGTTAGTCG
Km2	CAATAGCAGCCAGTCCCTTC
pQE3730F	CACCCATGGTATCACGGCGGATGTTGC
Km1	ACTGGGCTATCTGGACAAGG
<i>Gene cloning</i>	
Y3729-1	TAGCCCAACAGGAAGCGTAA
Y3729-XbaI-2	GATTCTAGATCGGCTTCATCGTGCATTAG
Y3731pBAD-EcoRI	GCCGAATTCGACTAACACAGAAAG
Y3731-down	GGCTCTGGGTCTATATCTGAG
Y3729-SD-fus-XbaI	ATCTCTAGATAGCCCAACAGGAAGCG
Y3729-R207-PstI	TTACTGCAGATCGTGCATTAGCCTTCTTACG
Y3730-SD-fus-XbaI	CAATCTAGACGTAAGAAGGCTAATGCACG
Y3730-G425-PstI	ATGCTGCAGATCCTAAACTTTCTGTGTTAGTCGTATTC
Y3730-R156-PstI	GTACTGCAGAACGCAATAAGCTGCCG
<i>Sequencing</i>	
hmsA_region_F	AGTGGCTCGCTTAGGTGG
hmsA_region_R	GCTTCGCCCTCAATCTCG
hmsA_region_Seq_F	CAACGAGCGGATGAAGCG
hmsA_region_Seq_R	GAGCCACGTTGAGTCTGC

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