

Supplementary material

The homologous components of flagellar type III protein apparatus **have** acquired a novel function to control twitching motility in a non-flagellated biocontrol bacterium

Alex M. Fulano¹, Danyu Shen¹, Miki Kinoshita², Shan-Ho Chou³, Guoliang Qian^{1*}

Table S1. Strains and plasmids used in this study

Strains and plasmids	Characteristics ^a	source
<i>Lysobacter enzymogenes</i>		
OH11	Wild type (CGMCC No. 1978), Km ^R	[1]
$\Delta flhA$	<i>flhA</i> in-frame deletion mutant, Km ^R	This study
$\Delta flhB$	<i>flhB</i> in-frame deletion mutant, Km ^R	This study
$\Delta flil$	<i>flil</i> in-frame deletion mutant, Km ^R	This study
$\Delta fliP$	<i>fliP</i> in-frame deletion mutant, Km ^R	This study
$\Delta fliQ$	<i>fliQ</i> in-frame deletion mutant, Km ^R	This study
$\Delta fliR$	<i>fliR</i> in-frame deletion mutant, Km ^R	This study
$\Delta pilA$	<i>pilA</i> in-frame deletion mutant, Km ^R	[2]
$\Delta pilB$	<i>pilB</i> in-frame deletion mutant, Km ^R	[2]
OH11(pBBR)	Wild type harboring plasmid pBBR1-MCS5, Gm ^R , Km ^R	This study
OH11(<i>pilA</i>)	Wild type harboring plasmid pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
$\Delta flhA(flhA)$	$\Delta flhA$ harboring pBBR- <i>flhA</i> , Gm ^R , Km ^R	This study
$\Delta flhB(flhB)$	$\Delta flhB$ harboring pBBR- <i>flhB</i> , Gm ^R , Km ^R	This study
$\Delta flil(flil)$	$\Delta flil$ harboring pBBR- <i>flil</i> , Gm ^R , Km ^R	This study
$\Delta fliR(fliR)$	$\Delta fliR$ harboring pBBR- <i>fliR</i> , Gm ^R , Km ^R	This study
$\Delta flhA(pilA)$	$\Delta flhA$ harboring pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
$\Delta flhB(pilA)$	$\Delta flhB$ harboring pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
$\Delta flil(pilA)$	$\Delta flil$ harboring pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
$\Delta fliR(pilA)$	$\Delta fliR$ harboring pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
$\Delta pilB(pilA)$	$\Delta pilB$ harboring pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
$\Delta pilQ(pilA)$	$\Delta pilQ$ harboring pBBR- <i>pilA</i> , Gm ^R , Km ^R	This study
<i>Escherichia coli</i> DH5 α	Host strain for molecular cloning	Laboratory collection
<i>Xanthomonas oryzae</i> PX099A	Source of the canonical FliC _{xoo}	Laboratory collection
<i>Salmonella</i>		
$\Delta flhA_{sal}$	<i>flhA</i> in-frame deletion mutant	[3]
$\Delta flhB_{sal}$	<i>flhB</i> in-frame deletion mutant	[3]
$\Delta flil_{sal}$	<i>flil</i> in-frame deletion mutant	[3]
$\Delta fliP_{sal}$	<i>fliP</i> in-frame deletion mutant	[3]
$\Delta fliQ_{sal}$	<i>fliQ</i> in-frame deletion mutant	[3]
$\Delta fliR_{sal}$	<i>fliR</i> in-frame deletion mutant	[3]

$\Delta fliA_{sal}$ (pTrc99A-FlhA)	$\Delta fliA_{sal}$ harboring pBBR- <i>fliA</i> , Amp ^R	This study
$\Delta fliB_{sal}$ (pTrc99A-FlhB)	$\Delta fliB_{sal}$ harboring pBBR- <i>fliB</i> , Amp ^R	This study
$\Delta fliI_{sal}$ (pTrc99A-FliI)	$\Delta fliI_{sal}$ harboring pBBR- <i>fliI</i> , Amp ^R	This study
$\Delta fliP_{sal}$ (pTrc99A-FliP)	$\Delta fliP_{sal}$ harboring pBBR- <i>fliR</i> , Amp ^R	This study
$\Delta fliQ_{sal}$ (pTrc99A-FliQ)	$\Delta fliQ_{sal}$ harboring pBBR- <i>fliQ</i> , Amp ^R	This study
$\Delta fliR_{sal}$ (pTrc99A-FliR)	$\Delta fliR_{sal}$ harboring pBBR- <i>fliR</i> , Amp ^R	This study
Plasmids		
pEX18GM	Suicide vector with a <i>sacB</i> gene, Gm ^R	[4]
pBBR1-MCS5	Broad-host-range vector with a Plac promoter, Gm ^R	[5]
pEX18- <i>fliA</i>	pEX18GM with two flanking fragments of <i>fliA</i> , Gm ^R	This study
pTrc99A-FlhA	An expression vector in <i>Salmonella</i> , Amp ^R	[3]
pEX18- <i>fliB</i>	pEX18GM with two flanking fragments of <i>fliB</i> , Gm ^R	This study
pEX18- <i>fliI</i>	pEX18GM with two flanking fragments of <i>fliI</i> , Gm ^R	This study
pEX18- <i>fliR</i>	pEX18GM with two flanking fragments of <i>fliR</i> , Gm ^R	This study
pEX18- <i>fliP</i>	pEX18GM with two flanking fragments of <i>fliP</i> , Gm ^R	This study
pEX18- <i>fliQ</i>	pEX18GM with two flanking fragments of <i>fliQ</i> , Gm ^R	This study
pBBR- <i>fliA</i>	pBBR1-MCS5 cloned with a 2385-bp fragment containing intact <i>fliA</i> and its predicted promoter, Gm ^R	This study
pBBR- <i>fliB</i>	pBBR1-MCS5 cloned with a 1557-bp fragment containing intact <i>fliB</i> and its predicted promoter, Gm ^R	This study
pBBR- <i>fliI</i>	pBBR1-MCS5 cloned with a 1562-bp fragment containing intact <i>fliI</i> , FLAG tag and its predicted promoter, Gm ^R	This study
pBBR- <i>fliR</i>	pBBR1-MCS5 cloned with a 1277-bp fragment containing intact <i>fliR</i> and its predicted promoter, Gm ^R	This study
pBBR- <i>pilA</i>	pBBR1-MCS5 cloned with a 737-bp fragment containing a PilA-Flag tag, Gm ^R	This study
pTrc99A-FlhA	pTrc99A cloned with the coding region of <i>fliA</i> of strain OH11, Amp ^R	This study
pTrc99A-FlhB	pTrc99A cloned with the coding region of <i>fliB</i> of strain OH11, Amp ^R	This study
pTrc99A-FliI	pTrc99A cloned with the coding region of <i>fliI</i> of strain OH11, Amp ^R	This study
pTrc99A-FliP	pTrc99A cloned with the coding region of <i>fliP</i> of strain OH11, Amp ^R	This study
pTrc99A-FliQ	pTrc99A cloned with the coding region of <i>fliQ</i> of strain OH11, Amp ^R	This study

^aKm^R, Gm^R and Amp^R: kanamycin-, gentamicin, and Ampicillin-resistance, respectively

Table S2. Primers used in this study

Primer	Sequence ^a (5'-3')	Purpose
In-frame deletion		
<i>flhA</i> -F1	CGGAATTCGATCGCCAAGAGCCAGGAAC (<i>EcoRI</i>)	To amplify a 794-bp upstream homologue arm of <i>flhA</i>
<i>flhA</i> -R1	GGGTTTGAAGGAGACGGGCGCGACGATGGTGGGGCAAGC	
<i>flhA</i> -F2	GCTTGCCCCACCATCGTCGCGCCCGTCTCCTTCAAACCC	To amplify a 799-bp downstream homologue arm of <i>flhA</i>
<i>flhA</i> -R2	GCTCTAGACTGTGCGATTGATGCGGCCGACGATGGTGGGGCAAGC (<i>XbaI</i>)	
<i>flhB</i> -F1	CCGGAATTCGTTCTGGTGTTCCTCGAC (<i>EcoRI</i>)	To amplify a 538-bp upstream homologue arm of <i>flhB</i>
<i>flhB</i> -R1	CGCCGACCCAGAAAAAGATCGGTCCTGCTGATTGGGCCGAAG	
<i>flhB</i> -F2	CTTCGGCCCAATGCAGGACCGATCTTTTCTGGGTCGGCG	To amplify a 464-bp downstream homologue arm of <i>flhB</i>
<i>flhB</i> -R2	GCTCTAGACCAGGCATCGGCGTTGAGGG (<i>XbaI</i>)	
<i>flip</i> -F1	CGGAATTCGGCTCACGGTCAAGGAACTG (<i>EcoRI</i>)	To amplify a 526-bp upstream homologue arm of <i>fliP</i>
<i>flip</i> -R1	GCGGCGAGGTCATTCATTTGCGATGGCGGACGAAGAAGGC	
<i>flip</i> -F2	GCCTTCTCGTCCGCCATCGCAAATGAATGACCTCGCCGC	To amplify a 645-bp downstream homologue arm of <i>fliP</i>
<i>flip</i> -R2	GCTCTAGAGGCAGCGAATGGAATAGTG (<i>XbaI</i>)	
<i>fliQ</i> -F1	CGGAATTCGAACATCCTGCTCTCGCTGG (<i>EcoRI</i>)	To amplify a 336-bp upstream homologue arm of <i>fliQ</i>
<i>fliQ</i> -R1	CCCAAGCTTGCTGTTTGGTCATTCGAG (<i>HindIII</i>)	
<i>fliQ</i> -F2	CCCAAGCTTCGCCGACCGCATCTTCCTCG (<i>HindIII</i>)	To amplify a 363-bp downstream homologue arm of <i>fliQ</i>
<i>fliQ</i> -R2	GCTCTAGACCCAGAACCATGCCGAAC (<i>XbaI</i>)	
<i>fliR</i> -F1	CGGAATTCACGAGGACGACACCGCGAC (<i>EcoRI</i>)	To amplify a 631-bp upstream homologue arm of <i>fliR</i>
<i>fliR</i> -R1	CCCAAGCTTCGGCGAGGCGATCAGGAAAC (<i>HindIII</i>)	
<i>fliR</i> -F2	CCCAAGCTTCCTTGGTGATGCTGACGCTC (<i>HindIII</i>)	To amplify a 923-bp downstream homologue arm of <i>fliR</i>
<i>fliR</i> -R2	GCTCTAGACGTAATGGGTGGGGTTGGTG (<i>XbaI</i>)	
<i>fliI</i> -F1	CCGGAATTCCTCTCCAGCGCCTCCAGCCT (<i>EcoRI</i>)	To amplify a 389-bp upstream homologue arm of <i>fliI</i>
<i>fliI</i> -R1	CCCAAGCTTCGGATGAACAAGCAACGCT (<i>HindIII</i>)	
<i>fliI</i> -F2	CCCAAGCTTCGACCACTTCGGCGGCGAG (<i>HindIII</i>)	To amplify a 457-bp downstream homologue arm of <i>fliI</i>
<i>fliI</i> -R2	CGGGGTACCCTGCGTGGTGTGCGGAAG (<i>KpnI</i>)	
Complementation and heterologous expression		
<i>flhA</i> -F	CCCAAGCTTCACCAACCCACCCATTACG (<i>HindIII</i>)	To amplify 2409-bp fragment containing the intact <i>flhA</i>
<i>flhA</i> -R	CGGAATTCCTCACTTATCGTCGTCATCCTTGTAATCGGCGGGC ACGGGTTTGA (<i>EcoRI</i>)	
<i>flhB</i> -F	CCCAAGCTTIGATGGACGCAATGTAGCCCC (<i>HindIII</i>)	To amplify 1331-bp fragment containing the intact <i>flhB</i>
<i>flhB</i> -R	CGGAATTCCTCACTTATCGTCGTCATCCTTGTAATCGCGCGGC TCGTCGTCGA (<i>EcoRI</i>)	
<i>fliI</i> -F	CCCAAGCTTCTGCGTGGTGTGCGGAAG (<i>HindIII</i>)	To amplify 1562-bp fragment containing the intact <i>fliI</i>
<i>fliI</i> -R	CGGAATTCCTCACTTATCGTCGTCATCCTTGTAATCTCCGAAC AACCTGCGCA (<i>EcoRI</i>)	
<i>fliP</i> -F	CCCAAGCTTGGCTCACGGTCAAGGAACTG (<i>HindIII</i>)	To amplify 1101-bp fragment containing the intact <i>fliP</i>
<i>fliP</i> -R	CGGAATTCCTCACTTATCGTCGTCATCCTTGTAATCTTGAAGC TGAGACCA (<i>EcoRI</i>)	
<i>fliQ</i> -F	CCCAAGCTTACCCTGACGGAACCTACCG (<i>HindIII</i>)	To amplify 676-bp fragment containing the intact <i>fliQ</i>
<i>fliQ</i> -R	CGGAATTCCTCACTTATCGTCGTCATCCTTGTAATCGTGCCGG ATCATGCCGG (<i>EcoRI</i>)	
<i>fliR</i> -F	CCCAAGCTTGAAGGACACGAGCAGGC (<i>HindIII</i>)	To amplify 1300-bp fragment containing the intact <i>fliR</i>
<i>fliR</i> -R	CGGAATTCCTCACTTATCGTCGTCATCCTTGTAATCTGGCGGC CCGCCGGGC (<i>EcoRI</i>)	
<i>pilA</i> -F	CCCAAGCTTCATCGCAAATCAACCCAC (<i>HindIII</i>)	To amplify 737-bp fragment containing the <i>pilA</i> -Flag fusion

<i>pilA-R</i>	CGGAATTCTTACTTATCGTCGTCATCCTTGTAATCGGTCGCA GCCTTGGTGC(<i>EcoRI</i>)	
<i>fliC</i> _{Xoo} F2	CCCAAGCTTATGTCGCTGAACGCTCAGCG (<i>HindIII</i>)	To amplify <i>fliC</i> _{Xoo} terminally fused with FLAG tag
<i>fliC</i> _{Xoo} F2	CGGAATTCTTACTTATCGTCGTCATCCTTGTAATCCTGCAGCA GGCTCAGCA(<i>EcoRI</i>)	

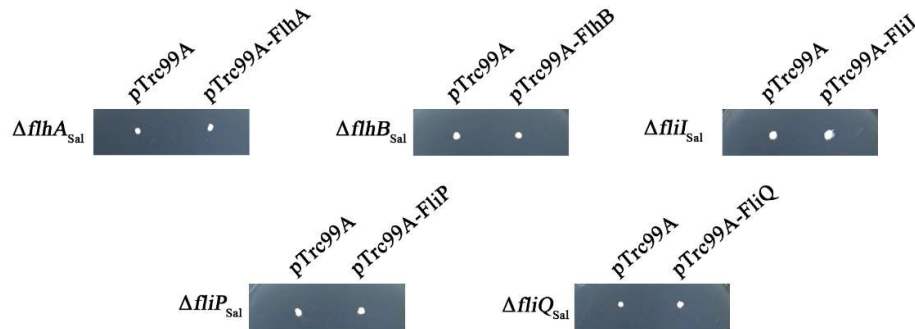


Figure S1. Cross-complementation of the *L. enzymogenes* FT3SS-like genes in the *Salmonella* mutants lacking the respective flagellar counterparts. The FlhA, FlhB, FliP, FliQ and FliI from strain OH11 failed to restore the swimming motility of the respective *Salmonella* mutants, indicated by $\Delta flhA_{sal}$, $\Delta flhB_{sal}$, $\Delta fliP_{sal}$, $\Delta fliQ_{sal}$ and $\Delta fliI_{sal}$, respectively. The photograph was taken after incubation of the transformed strain at 30°C for 24 hours.

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