

Supplementary Material

Supplementary Table S1. Strains, plasmids and primers used in this study

Strains/Plasmids/Primers	Description	Source
Strains		
<i>E. coli</i> CA434	Donor cells for conjugation transformation	[34]
<i>Ct</i>	ATCC 25755	[27]
pcloneEZ- <i>Plac</i> -repeat	Used for <i>Plac</i> promoter amplification	Stored in laboratory
pcloneEZ-repeat-	Used for terminator amplification	Stored in laboratory
<i>Ct</i> /Δ <i>abrB</i>	ATCC 25755 with pM3/Δ <i>abrB</i>	This study
<i>Ct</i> /Δ <i>abrB</i>	ATCC 25755 with pM2/Δ <i>abrB</i>	This study
<i>Ct</i> /Δ <i>abrB</i> -Δ <i>abrB</i>	<i>Ct</i> /Δ <i>abrB</i> with pM2/Δ <i>abrB</i>	This study
<i>Ct</i> /Δ <i>spo0A</i>	ATCC 25755 with pM3/Δ <i>spo0A</i>	This study
<i>Ct</i> /Δ <i>sigF</i>	ATCC 25755 with pM3/Δ <i>sigF</i>	This study
<i>Ct</i> /Δ <i>sigE</i>	ATCC 25755 with pM3/Δ <i>sigE</i>	This study
<i>Ct</i> /Δ <i>sigG</i>	ATCC 25755 with pM3/Δ <i>sigG</i>	This study
<i>Ct</i> /Δ <i>sigK</i>	ATCC 25755 with pM3/Δ <i>sigK</i>	This study
Plasmids		
pMTL82151	ColE1 ori; Cm ^R ; pBP1 ori; <i>TarJ</i>	[63]
pMTL82151-Em	ColE1 ori; Em ^R ; pBP1 ori; <i>TarJ</i>	Stored in laboratory
pMTL83151	ColE1 ori; Cm ^R ; pBP1 ori; <i>TarJ</i>	Stored in laboratory
pM3/Δ <i>abrB</i>	From pMTL83151; <i>Plac</i> -Δ <i>abrB</i>	This study
pM2/Δ <i>abrB</i>	From pMTL82151-Em; <i>Pcat1</i> -Δ <i>abrB</i>	This study
pM3/Δ <i>spo0A</i>	From pMTL83151; <i>Plac</i> -Δ <i>spo0A</i>	This study
pM3/Δ <i>sigF</i>	From pMTL83151; <i>Plac</i> -Δ <i>sigF</i>	This study
pM3/Δ <i>sigE</i>	From pMTL83151; <i>Plac</i> -Δ <i>sigE</i>	This study
pM3/Δ <i>sigG</i>	From pMTL83151; <i>Plac</i> -Δ <i>sigG</i>	This study
pM3/Δ <i>sigK</i>	From pMTL83151; <i>Plac</i> -Δ <i>sigK</i>	This study
Primers		
<i>Pcat1</i> -F	gaaacagctatgaccgcgccgcGTAGACTTTAAGGATGGAACCTTTGA	
Pro-F	TGAAGTACATCACCGACGAGCAAG	
Pro-R	TGCTGCAAGGCGATTAAGTTGGGT	
<i>Plac</i> -F	gaaacagctatgaccGCGGCCGCTTATATACTTGTTTATTTACTTGATTAT	
<i>Plac</i> -repeat(<i>abrB</i>)-s-R	agatacatgctggttcataatttcttaataATTTAAATACATCTCATGTAAAGGTTT	
spacer(<i>abrB</i>)-pb-F	aatatgaaccagcatgtatcttctgGTTGAACCTTAACATGAGATGTATT	
spacer(<i>abrB</i>)-term-R	gaaatataggagctcATAAAAAAATTGTAGATAAACTATTTTATA	
<i>abrB</i> -H1-F	ctacaattttttatgagctcCTATATTTCCATCATATATAAAATCTGC	
<i>abrB</i> -H1-R	gtttttacagaagaaggtaccTTATTTTCCTCCCTAAGTTCCA	

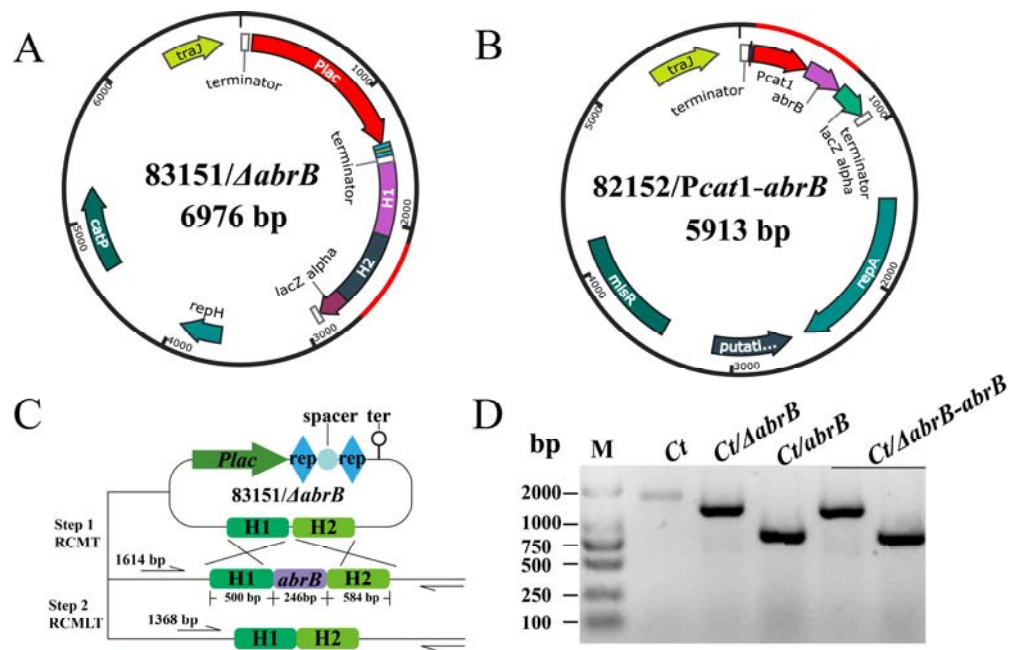
Supplementary Table S1. Cont.

Strains/Plasmids/Primers	Description	Source
<i>abrB</i> -H2-F	ggaaaataaggtaccTTCTTCTGTAAAAACAGGCTATTT	
<i>abrB</i> -H2-R	tctccatggacgcgtgacgtcCGGCACCTAGAATTCAATCA	
<i>abrB</i> -conf-F	AAGCTCTCCTTAGATACTGATTTAATTATTT	
<i>abrB</i> -conf-R	CTACACAAACTAAATTAATAGCAGATTTAGAC	
<i>Pcat1</i> (<i>abrB</i>)-R	tcttacaacacctgttgatttcatAAAAACCACCCTTTCATAAATTA	
<i>abrB</i> -F	aaagggtgggttttgcggccgcATGAAATCAACAGGTGTTGTAAGA	
<i>abrB</i> -R	tctccatggacgcgtgacgtcTTATCTTCCTTCTTTTAATTCATTTA	
Plac-repeat(<i>spo0A</i>)-s-R	taatagtagtgagagcaagaaacgggtatATTTAAATACATCTCATGTTAAGGTTTC	
spacer(<i>spo0A</i>)-pb-F	ttcttgctctcactactattagctatatcaGTTGAACCTTAACATGAGATGTATTAA	
spacer(<i>spo0A</i>)-term-R	gcttctaagtcctatgagctcATAAAAAAATTGTAGATAAAACTATTTTATAAA	
<i>spo0A</i> -H1-F	ttttttatgagctcATAGGACTTAGAAGCGAAATAAAAGAAG	
<i>spo0A</i> -H1-R	attgatatactttatctagaATATTTTACTCCCCTTTTATTAATAAATTCC	
<i>spo0A</i> -H2-F	gtaaaatattctagaTAAAGTGATATCAATGGTTAACAGGTTT	
<i>spo0A</i> -H2-R	acgacggccagtgccaagcttGTTATCTGTGATTTCCGGTATTCTATTATTC	
<i>spo0A</i> -conf-F	CCCAGTGC GGAGTATTTGGTA	
<i>spo0A</i> -conf-R	ATTATCGTGTAGTCCCTAAATTCGTG	
Plac-repeat(<i>sigF</i>)-s-R	tctatcaagagaactgggtctccatcatccATTTAAATACATCTCATGTTAAGGTTTC	
spacer(<i>sigF</i>)-pb-F	agcaccagttctcttgatagataagGTTGAACCTTAACATGAGATGTATT	
spacer(<i>sigF</i>)-term-R	gtcatacaccgagctcATAAAAAAATTGTAGATAAAACTATTTTATA	
<i>sigF</i> -H1-F	ctacaattttttatgagctcGGTGTATGACAGCGTTGAGC	
<i>sigF</i> -H1-R	acgaaacctacttatggatccATTATTCACCTAACTTAATGATTTAAAA	
<i>sigF</i> -H2-F	gggtgaataatggatccATAAGTAGGTTTCGTAACATAAAAAATAT	
<i>sigF</i> -H2-R	tctccatggacgcgtgacgtcGCCTTCTGGAGTTGTAAGTGA	
<i>sigF</i> -conf-F	GGATAGTTCAGGCATAGGTGTT	
<i>sigF</i> -conf-R	TTAACTCCCTACTACTCCGCTT	
Plac-repeat(<i>sigE</i>)-s-R	tgtaccactgatatacaaatcttctacatATTTAAATACATCTCATGTTAAGGTTTC	
spacer(<i>sigE</i>)-pb-F	atttgatatcagtggtgacaataggGTTGAACCTTAACATGAGATGTATT	
spacer(<i>sigE</i>)-term-R	cctaagagccggtaccATAAAAAAATTGTAGATAAAACTATTTTATA	
<i>sigE</i> -H1-F	ctacaattttttatgggtaccGGCTCTTAGGCTCTATTATGATT	
<i>sigE</i> -H1-R	atactaaagacatgcggatccAATTTTCCTCCAATTAATCATAGT	
<i>sigE</i> -H2-F	aggaaaattggatccGCATGTCTTTAGTATAAAATATAGCAT	
<i>sigE</i> -H2-R	tctccatggacgcgtgacgtcCTCTAGCTTGAAGTGCCTTATAA	
<i>sigE</i> -conf-F	TTGCATATTTATAATGTATTCTATGCT	
<i>sigE</i> -conf-R	AATGATACTGGATCTTGAATTGC	
Plac-repeat(<i>sigG</i>)-s-R	attgctttataagaccaacacatccgaccATTTAAATACATCTCATGTTAAGGTTTC	
spacer(<i>sigG</i>)-pb-F	tggttggtcttataaaagcaatagatGTTGAACCTTAACATGAGATGTATT	

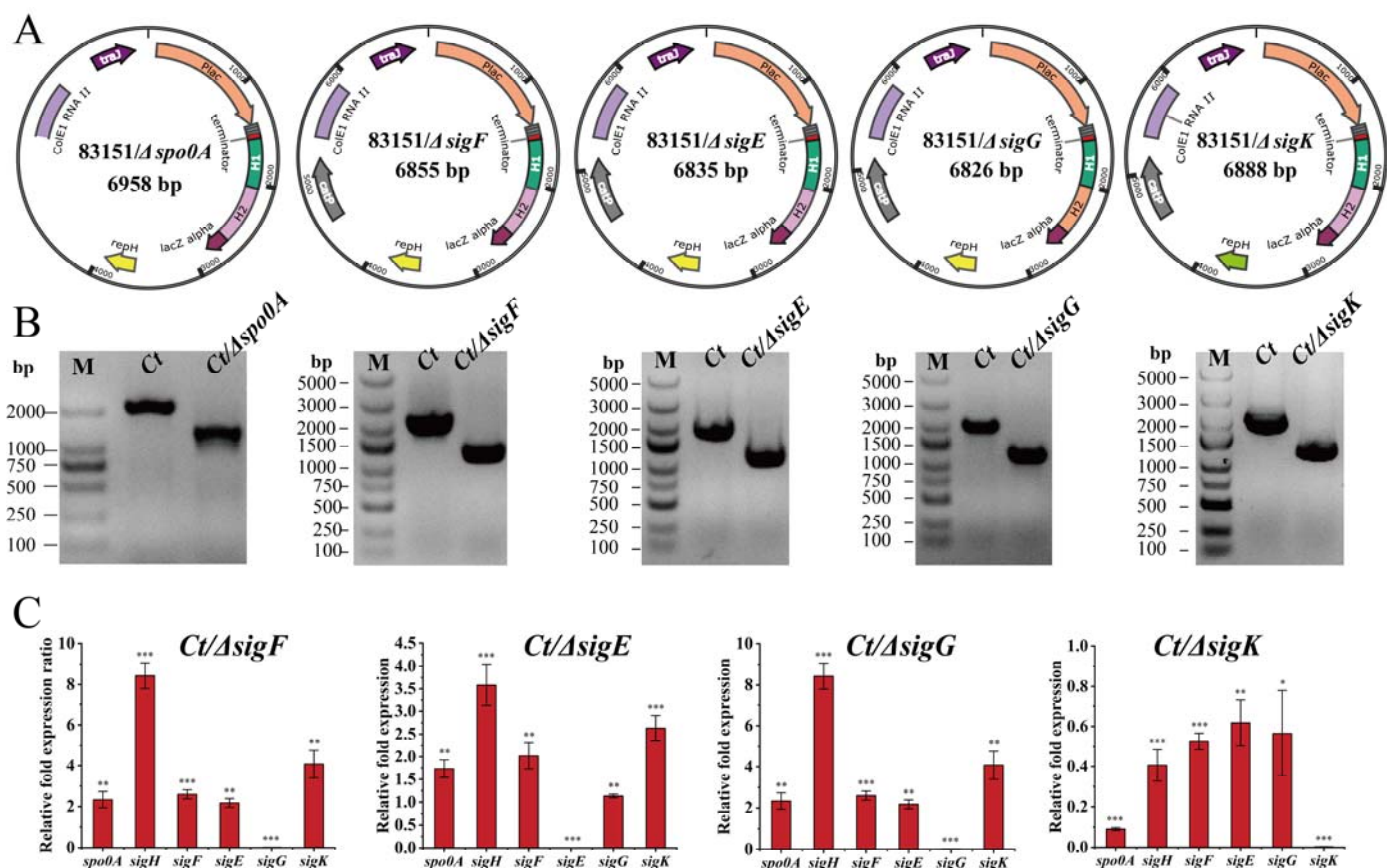
Supplementary Table S1. *Cont.*

Strains/Plasmids/Primers	Description	Source
spacer(<i>sigG</i>)-term-R	tatgttgccgagctcATAAAAAAATTGTAGATAAACTATTTTATA	
<i>sigG</i> -H1-F	ctacaattttttatgagctcGGCAACATATGCATCCAGA	
<i>sigG</i> -H1-R	tacaatataaatataggtaccGCAATCAGCCCCTTTAGAA	
<i>sigG</i> -H2-F	gctgattgcggtaccTATATTTATATTGTAAGTAATATTAATGGG	
<i>sigG</i> -H2-R	tctccatggacgcgtgacgtcGGGCATTTCACTTCTCTTCA	
<i>sigG</i> -conf-F	TACTGGAGTCAATGTAGAAGATTTG	
<i>sigG</i> -conf-R	ATAACTAATATAGGCAAGTCTTCAATT	
Plac-repeat(<i>sigK</i>)-s-R	ttccccggataagaatattttttacaatgATTAAATACATCTCATGTTAAGGTTC	
spacer(<i>sigK</i>)-pb-F	aaaatattcttatccgggaaggatGTTGAACCTTAACATGAGATGTATT	
spacer(<i>sigK</i>)-term-R	tcactactggagctcATAAAAAAATTGTAGATAAACTATTTTATA	
<i>sigK</i> -H1-F	ctacaattttttatgagctcCAGTAGTGAATTCAGGTATATATGTAAAA	
<i>sigK</i> -H1-R	ttaatcttaagaataggtaccAGCTGCTCCTCCTAATAGTGC	
<i>sigK</i> -H2-F	ggagcagctggtaccTATTCTTAAGATTAAAAAATATATAAAAATTT	
<i>sigK</i> -H2-R	tctccatggacgcgtgacgtcCCACATCCAGTTGGTCCTG	
<i>sigK</i> -conf-F	TTGCATATTTATAATGTATTCTATGCT	
<i>sigK</i> -conf-R	TGTATCTTTACCTATTTCTCTTTGATT	

Note: Small letters are homologous sequences and capital letters are specific sequences. Underline marks are enzyme digestion sites



Supplementary Figure S1 Construction and verification of the *Ct/ΔabrB*, *Ct/abrB* and *Ct/ΔabrB-abrB* strains. (A) The *abrB* knockout plasmid; (B) The *abrB* overexpression plasmid; (C) Schematic illustrating the work flow of *abrB* gene deletion using the lactose inducible CRISPR-Cas system. (D) Gel electrophoresis verification of *abrB* mutant. M: Marker.



Supplementary Figure S2 Construction of *Ct/Δspo0A* and *Ct/Δsigma* strains. (A) Construction of the knockout plasmid of *spo0A* and sigma factors; (B) Gel electrophoresis verification of *spo0A* and sigma factors mutant; (C) qRT-PCR analysis

of genes related to sporulation transcription levels at 72 h. (ns, non-significant; *, $p<0.05$; **, $p<0.01$; ***, $p<0.001$, t-test).