

SpoVG Regulates Cell Wall Metabolism and Oxacillin Resistance in
Methicillin-Resistant *Staphylococcus aureus* Strain N315

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Running title: SpoVG Regulates Oxacillin Resistance

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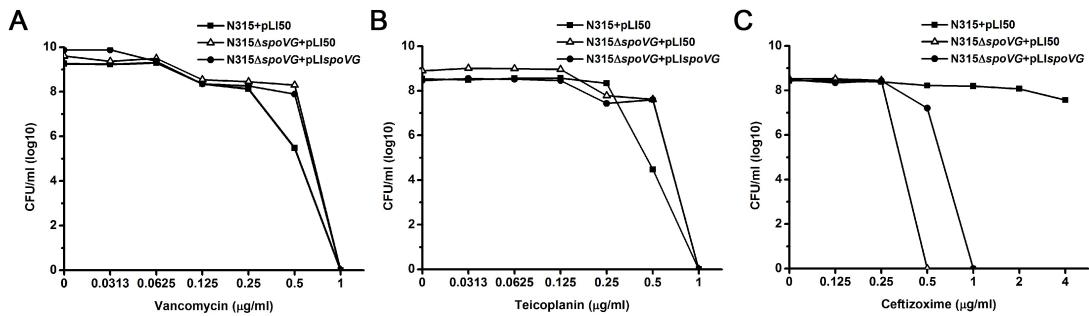


Figure S1. Antibiotic susceptibility assay. The antibiotic resistance levels were evaluated by population analysis profiles as recommended by the Clinical and Laboratory Standards Institute. The colonies were counted after incubation at 37°C for 24 hours. (A) and (B) There is no dramatic difference in vancomycin and teicoplanin resistance between the *spoVG* mutant (N315 Δ *spoVG*+pLI50) and wide-type (N315+pLI50) strains. (C) Deletion of *spoVG* reduced ceftizoxime resistance in strain N315 and the complementary (N315 Δ *spoVG*+pLIs*spoVG*) strain has a partial effect.

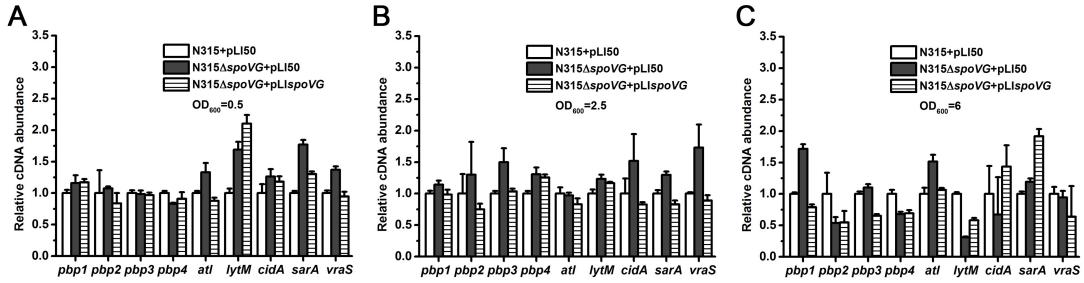


Figure S2. Transcriptional analysis of cell wall-related genes. Comparison of the relative transcription levels of several cell wall biosynthesis- and hydrolysis-related genes in the wild-type (N315+pLI50), the *spoVG* mutant (N315Δ*spoVG*+pLI50), and the *spoVG* complementary (N315Δ*spoVG*+pLIs*spoVG*) strains. (A), (B), and (C) The relative transcription levels of *pbp1*, *pbp2*, *pbp3*, *pbp4*, *atl*, *lytM*, *cida*, *sarA*, and *vraS* genes in early exponential ($OD_{600}=0.5$), mid-logarithmic ($OD_{600}=2.5$) and early stationary ($OD_{600}=6$) phases were not significantly affected in the *spoVG* mutant (fold change <2) except for *lytM* of early stationary phase.

Table S1. Primers used in this study

Primer name	Oligonucleotide ^a (5'-3')	Application
<i>spoVG</i> -mutant-u	GCGggtaccAATTGCCAGTATTACATGG	pBTs Δ <i>spoVG</i> ,
p-F		pLI <i>spoVG</i>
<i>spoVG</i> -mutant-u	<u>GCAAATGTATTACATCGCTACCCCCCTAT</u>	pBTs Δ <i>spoVG</i>
p-R	AGTATATATCTC	
<i>spoVG</i> -mutant-d	TAGCGATGTAATAACATTGC	pBTs Δ <i>spoVG</i>
own-F		
<i>spoVG</i> -mutant-d	GCGgtcgacTATTCACCTTGCGCATTATC	pBTs Δ <i>spoVG</i>
own-R		
P <i>spoVG</i> -R	ACTAAA ACTCCTTTATGAAAAC	pLI <i>spoVG</i>
<i>spoVG</i> -complem	<u>TTCATAAAAGGAGTTTAGTATGAAAGT</u>	pLI <i>spoVG</i>
ent-F	GACAGATGTAAGACT	
<i>spoVG</i> -complem	GCGggtaccTGCCGACAAAGCTTATGCT	pLI <i>spoVG</i>
ent-R		
SpoVG-exp-F	CATGccatgggAATGAAAGTGACAGATGTA AGAC	pETSpoVG
SpoVG-exp-R	CCGctcgagAGCTTCTTCTGAATCTTCTGAT GTAGC	pETSpoVG
RTQ- <i>hu</i> -F	AAAAAGAAGCTGGTCAGCAGTAG	qRT-PCR
RTQ- <i>hu</i> -R	TTTACGTGCAGCACGTTCAC	qRT-PCR

Table S1-continued

Primer name	Oligonucleotide ^a (5'-3')	Application
RTQ- <i>lytN</i> -F	TGACACCATTAGTAGAACCA	qRT-PCR
RTQ- <i>lytN</i> -R	CATTGCCATCCATAACTACC	qRT-PCR
RTQ- <i>lrgA</i> -F	TTATGCCAATTCTATGC	qRT-PCR
RTQ- <i>lrgA</i> -R	CGAAGAGTAAGCCAATG	qRT-PCR
RTQ- <i>arlS</i> -F	CGAATGATAACACAGTGAGAG	qRT-PCR
RTQ- <i>arlS</i> -R	CTTGC GTTGT AATTGGTTCT	qRT-PCR
RTQ- <i>lytS</i> -F	GGCACGAGAGTTACTATTAGA	qRT-PCR
RTQ- <i>lytS</i> -R	GCACGC ACTTGACTTAAC	qRT-PCR
RTQ- <i>sleI</i> -F	CACCA GTATT CAGTCACCAA	qRT-PCR
RTQ- <i>sleI</i> -R	CAGTTATTAGCATTCCACCAAT	qRT-PCR
RTQ- <i>mecA</i> -F	CGACTTCACATCTATTAGGTTAT	qRT-PCR
RTQ- <i>mecA</i> -R	ACACGATAGCCATCTTCA	qRT-PCR
RTQ- <i>femA</i> -F	ACACA ACAAG CGAGATAAC	qRT-PCR
RTQ- <i>femA</i> -R	CAGCAGAGATA GGTAATT CATT	qRT-PCR
RTQ- <i>blaZ</i> -F	TTTCCATTGCGATAAGTTT	qRT-PCR
RTQ- <i>blaZ</i> -R	TCCAGTTAGATATGAGATAGAAT	qRT-PCR
RTQ- <i>lytM</i> -F	CATT CGTAGAT GCTCAAGGA	qRT-PCR
RTQ- <i>lytM</i> -R	CTCGCTGTGTAGTCATTGT	qRT-PCR
RTQ- <i>cida</i> -F	GCAGTCATTATCATAGGAA	qRT-PCR
RTQ- <i>cida</i> -R	CATT CATAAGCGTCTACA	qRT-PCR

Table S1-continued

Primer name	Oligonucleotide ^a (5'-3')	Application
RTQ- <i>atl</i> -F	AGCACCAACGGATTAC	qRT-PCR
RTQ- <i>atl</i> -R	CATACTCAGCACTGTCT	qRT-PCR
RTQ- <i>sarA</i> -F	GTGCATTAACAAGAATTAAATACAGT	qRT-PCR
RTQ- <i>sarA</i> -R	TTGACATACATCAGCGAAAAC	qRT-PCR
RTQ- <i>pbp2</i> -F	ATGGTGTATCTGGAGTAA	qRT-PCR
RTQ- <i>pbp2</i> -R	CATAGTGTATTGAGGTGTAA	qRT-PCR
RTQ- <i>pbp4</i> -F	TATGAACAATAAGTGCTAACCC	qRT-PCR
RTQ- <i>pbp4</i> -R	GACCTCCAACGTAGAAG	qRT-PCR
RTQ- <i>vraS</i> -F	TGGTCTAATGGTGGTTCT	qRT-PCR
RTQ- <i>vraS</i> -R	CTTCACGATTCTGTTAGTCA	qRT-PCR
<i>plytN</i> -F	AGTTAAAAGATGATATAAAAT	EMSA
<i>plytN</i> -R-biotin	AGCGTACTTTACTTTGTTGT	EMSA
<i>plytN</i> -R	AGCGTACTTTACTTTGTTGT	EMSA
<i>pfemA</i> -F	TAATATGAGCAAACCCAAACG	EMSA
<i>pfemA</i> -R-biotin	AGGCACCAAACTCTTAGC	EMSA
<i>pfemA</i> -R	AGGCACCAAACTCTTAGC	EMSA
<i>plytSR</i> -F	GATTAGAAACTGTTGGCAAAG	EMSA
<i>plytSR</i> -R-biotin	TTGTTAGCGATAGCACGATA	EMSA
<i>plytSR</i> -R	TTGTTAGCGATAGCACGATA	EMSA

Table S1-continued

Primer name	Oligonucleotide ^a (5'-3')	Application
RTQ- <i>pbp1</i> -F	AGTCAGCGACCAACATT	qRT-PCR
RTQ- <i>pbp1</i> -R	AAGCACCTTCTTGAATAGC	qRT-PCR
RTQ- <i>pbp3</i> -F	TTCAGCCGTATCCAACATT	qRT-PCR
RTQ- <i>pbp3</i> -R	GTGCCAAGAGGTCGTATT	qRT-PCR

^a Lowercase letters indicate restriction sites. Underlined letters indicate overlap sequences used for SLiCE.