

**S3 Table. Oligonucleotides used in this study**

Oligonucleotide <sup>a</sup>	Sequence (5' → 3') <sup>b</sup>	Use
RTcabA	CAGATTGTCGCTCATGCGG	RT-PCR of <i>cabABC</i>
RTcabC2	AGATAGCTGATTGGCGTCCGTG	
cabAexp_F	<u>CCATGGCTGTTATTCTGGAAC</u> TG	Amplification of the <i>cabA</i> coding region
cabAexp_R	<u>CTCGAGGAATTGGAACATGTCATC</u>	
dcpA_F1	ACCCCGCTTGTGAGGTGAAC	
dcpA_R1	<u>GATTCCATGGCGTTAGCCA</u> ACTGTAAT	
dcpA_F2	<u>AACGCCATGGAATCATATTAGGTAA</u> AGGAAG	Integration of P <sub>BAD</sub> with <i>dcpA</i>
dcpA_R2	GAGCCC GTTAGATCAGAGACATGTCAG	
pBAD24_F	<u>CCATGGTGCATAATGTGCCTGTC</u>	
pBAD24_R	CCGGGTACCATGGTGAATTCT	
cabA_F1	<u>ACTAGTGGTGGAGCGAAGAAGGAA</u>	
cabA_R1	<u>CGGGATCCCGTTCCAGAATAAAC</u> CA	Construction of the <i>cabA</i> mutant
cabA_F2	<u>ACGGATCCC GGATGACATGTT</u>	
cabA_R2	<u>GCATGCGATAAAAGAGAGGAACCAC</u>	
cabB_F1	GAAGTTGATTGGTGTGACAGAACAG	
cabB_R1	<u>AAGGATCCAAGCGTCTGTAAATCTCTC</u>	Construction of the <i>cabB</i> mutant
cabB_F2	<u>TTGGATCCTTGCTCATATCCGAG</u>	
cabB_R2	GACGGCATGGACAAACCGTC	
cabC_F1	AATAACGGAGAGAAATGGTGAGACGTTAG	
cabC_R1	<u>ATGGATCCTTCCCAGACAATCTCACT</u>	Construction of the <i>cabC</i> mutant
cabC_F2	<u>AAGGATCCATGACCAAGATGCTCAG</u>	
cabC_R2	GTACGTTTCGACATCCGATATGATGAC	
brpA_F1	GCGGGTTTAC GCCTATCGTGG	
brpA_R1	<u>CATCGGATCCATGTGCACATCTCGT</u>	Construction of the <i>brpA</i> mutant
brpA_F2	<u>ACATGGATCCGATGCAGCTCATTG</u>	
brpA_R2	GCAACTGTTGTTGATATT CGCGAGGC	
cabAcom_F	<u>AGTCCATGGTCCAATAATGGAATCAAAGAGG</u>	Complementation of the <i>cabA</i> mutant
cabAcom_R	<u>CTGCATGCCACCTAGAAGTAGTG</u>	
cabBcom_F	<u>AGCCATGGTAGTAATT TGCAAAATT</u> C	Complementation of the <i>cabB</i> mutant
cabBcom_R	<u>AAGCATGCAACGTCTACCATTTC</u>	
cabCcom_F	<u>AGCCATGGCTAATCAAATAACGGAG</u>	Complementation of the <i>cabC</i> mutant
cabCcom_R	<u>GAGCATGCATACATCAGTAATCATTG</u>	
qRTcabA_F	TTGGTTGCTGGCTCTGGTGAC	
qRTcabA_R	ACTGTCTATACGCACTGTGTCCTC	qRT-PCR
qRTcabB_F	GCCATTGCCAGACCCAGAG	

qRTcabB_R	CCGATAATACCAACCGCACACC
qRTcabC_F	<u>TTGGCGGTGGTATTGGCTACTG</u>
qRTcabC_R	<u>TGTTGAATTGCCTGGCGTTGAC</u>

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<sup>a</sup> The oligonucleotides were designed using the *V. vulnificus* CMCP6 genome sequence.

<sup>b</sup> Regions of oligonucleotides not complementary to corresponding genes are underlined.