

Supplemental Material for

The TonB3 system in the human pathogen *Vibrio vulnificus* is under the control of the global regulators Lrp and CRP.

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TABLE S1. Primers used in this work

Primer name	Sequence (5' to 3')	Use
42-100	CTGCAGTCTCATGTATATCTGTCGCGAAC	<i>lacZ</i> fusion
42-300	CTGCAGTCACTACAATAGATCCACCCCTCCA	<i>lacZ</i> fusion/EMSA/ footprinting
42-400	CTGCAGATTGATAACCACCAGACTCATCTT	<i>lacZ</i> fusion
42-498	CTGCAGACTACATAAAAATGCAATTGCAC	<i>lacZ</i> fusion/EMSA/ footprinting
42-632	CTGCAGAAGTATTGTGATGAAAGCATGAC	<i>lacZ</i> fusion/EMSA/ footprinting
42Pst	CTGCAGCTGAACAGCTTCAATCTCTGG	<i>lacZ</i> fusion
42Bam	GGATCCTGCGAAAGCTATCGGGGACAT	<i>lacZ</i> fusion/EMSA/ footprinting
42-548	GCTGCAGCCAATCACAGAATATCAAACGT	<i>lacZ</i> fusion
42-300Rev	TGTTGGAGGGTGGATCTATTGTAG	EMSA/ footprinting
42-dintREV	GTTGGAGGGTGGATCTATTGTAGTCTCATTGGC TGAAGCTTAAAC	deletion
42-dintFOR	GTTTAAAGCTTCAGCCAAATGAGACTACAATAGA TCCACCCCTCCAAC	deletion
42-477Bam	CGGATCCGTTAGAATGGCTGACTGTTATTG	EMSA/ footprinting
42-206-For	CTGCAGGCCAAATCACAGATAATCATGCTG	EMSA
42-206-Rev	CAGCATGATTATCTGTGATTGGC	EMSA
M13F	ACGTTGTAAAACGACGGCCAG	footprinting
M13R	CAGGAAACAGCTATGAC	footprinting

delta632-498-1	CGCGTGCACATGACGTCTTCG	deletion
delta632-498-3	CTGTTAGAATGGCTGACTGTTATTGTCATGCTT CATCACAAATACTT	deletion
delta632-498-4	AAGTATTGTGATGAAAGCATGACAATAAACAGT CAGCCATTCTAACAG	deletion
delta632-498-2	TTTGC CGAGTGTACCACCTGTG	deletion
Lrp-1	AACAGCCATACGAGCAGCGTTAG	deletion
Lrp-2	AAACAGCAACACCAACCGTTGTAC	deletion
Lrp-3	ACGTGTTTAATCACAAGTTGGTTCGGCTTTGT AGTTATCTACCAT	deletion
Lrp-4	ATGGTAGATAACTACAAAAAGCCAACCAACTT GTGATTAAAACACGT	deletion
CRP-1	GATGTGCTGTTTATGAAGGACTG	deletion
CRP-2	TAAAAGTGGATGAATGCTCGGAGG	deletion
CRP-3	CGAGTACCGTAAACAACGATAGTCGGTTGAGGT TTACCTAGAACCAT	deletion
CRP-4	ATGGTTCTAGGTAAACCTCAAACCGACTATCGTT GTTTACGGTACTCG	deletion
Lrp-revSAC	GGAGCTCTGCCGCTCGGATTCTAAAGTC	Complementat ion/pET200 cloning
Lrp-ForPET	CACCATGGTAGATAACTACAAAAAGCCG	pET200 cloning
Lrp-ForPST	GCTGCAGCTATGTGTAGGGTGTAAATGTCG	complemen- tation
CRP-pETFOR	CACCATGGTTCTAGGTAAACCTCAAACC	pET200 cloning
CRP-REVSAC	GGAGCTCGTTGCTGCTGAAGCGTAGTGCTC	complemen- tation/pET200 cloning
CRP-	TTCTAGACCGTGATCATGTGCACAGTTTGCG	complemen-

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FORXBA		tation
TONB3F-qRT	TCTTGAGTTACCGTCAACG	qRT-PCR
TONB3R-qRT	GTTTGCCATCGACCATCTG	qRT-PCR
Lrp-ForqRT	AATGAATTGCAGAAAGATGGTCG	qRT-PCR
Lrp-RevqRT	ACATGGAGTTGGAGAAAGCC	qRT-PCR
CRP-qRTFOR	GGTGACTTAGCATTCTAGACG	qRT-PCR
CRP-qRTREV	TGACCGATTCTTGACGAGTG	qRT-PCR
23S-Forward	AGGCATGCTGGAGGTATCAGAA	qRT-PCR
23S-Reverse	AACGTTGGACAGGAACCCTTG	qRT-PCR

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