BfvR, an AraC-family regulator, controls biofilm formation and pH6 antigen production in opposite ways in *Yersinia pestis* biovar Microtus

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Short running title: BfvR controls biofilm and virulence

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Abstract

Biofilm formation is critical for blocking flea foregut and hence for transmission of Y. pestis by

flea biting. In this study, we identified the regulatory role of the AraC-family transcriptional

regulator BfvR (YPO1737 in strain CO92) in biofilm formation and virulence of Yersinia pestis

biovar Microtus. Crystal violet staining, Caenorhabditis elegans biofilm assay, colony

morphology assay, intracellular c-di-GMP concentration determination, and BALB/c mice

challenge were employed to reveal that BfvR enhanced Y. pestis biofilm formation while

repressed its virulence in mice. Further molecular biological assays demonstrated that BfvR

directly stimulated the expression of hmsHFRS, waaAE-coaD, and hmsCDE, which, in turn,

affected the production of exopolysaccharide, LPS and c-di-GMP, respectively. In addition, BfvR

directly and indirectly repressed psaABC and psaEF transcription, respectively. We concluded that

the modulation of biofilm- and virulence-related genes by BfvR led to increased biofilm formation

and reduced virulence of Y. pestis biovar Microtus.

Keywords: Yersinia pestis; BfvR; biofilm; virulence; transcriptional regulation.

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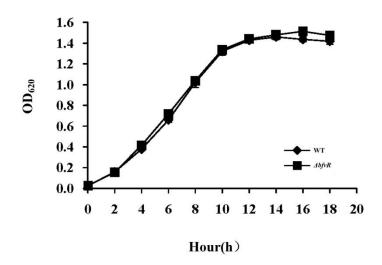


FIGURE S1. Bacterial growth curves. Overnight cell cultures at an OD620 value of approximately 1.0 were diluted 1:20 into 18 ml of LB medium with a pH value of 6.0. Bacteria were then grown at 26 °C with shaking at 230 rpm, and the OD620 values were monitored at 3-h intervals until the cell cultures reached the stationary growth phase. Three individual cultures (i.e., three biological replicates) were determined for each strain.

OD 620: Optical density at 620 nm; WT: Wild-type.

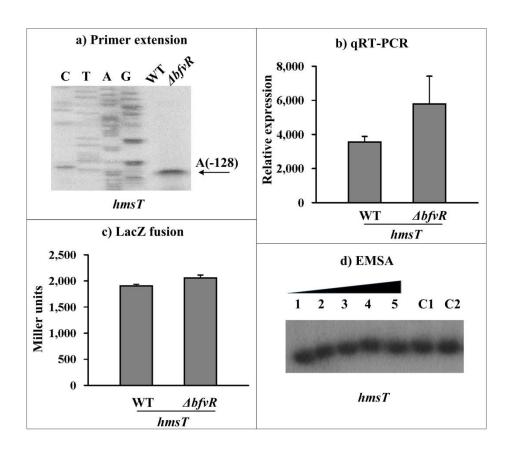


FIGURE S2. Regulation of *hmsT* **by BfvR.** See FIGURE 2 for the annotations of primer extension (**a**), quantitative RT-PCR (**b**), LacZ fusion (**c**), and EMSA (**d**). The promoter-proximal region was amplified from -410 to +42.

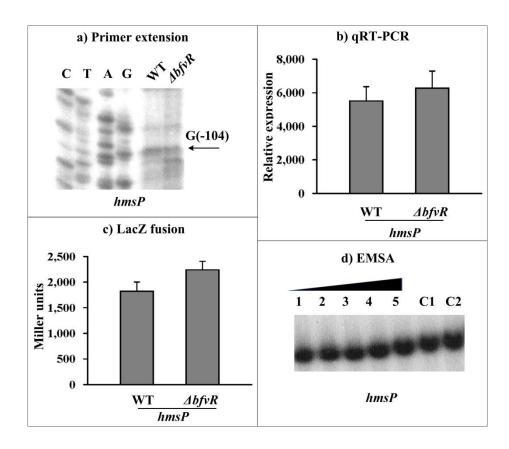


FIGURE S3. Regulation of *hmsP* **by BfvR.** See FIGURE 2 for the annotations of primer extension (**a**), quantitative RT-PCR (**b**), LacZ fusion (**c**), and EMSA (**d**). The promoter-proximal region was amplified from -659 to +65.

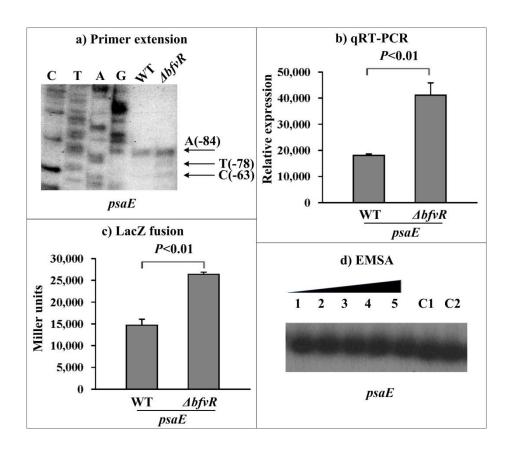


FIGURE S4. Regulation of *psaE* **by BfvR.** See FIGURE 2 for the annotations of primer extension (**a**), quantitative RT-PCR (**b**), LacZ fusion (**c**), and EMSA (**d**). The promoter-proximal region was amplified from -563 to +93.