

Table S3. Plasmids used in this study

Plasmid name	Description	Source/ Reference
pMYSH6504	pBR322 derivative carrying <i>S. flexneri</i> <i>virF</i> gene	(1)
pF-M4G	pMYSH6504 derivative with ATG to GGG mutation at positions 71-73 in <i>virF</i> (primer pairs used: ML-296 /ML-297)	This study
pF-M81L	pMYSH6504 derivative with ATG to CTG mutation at positions 311-313 in <i>virF</i> (ML-394/ML-395)	This study
pF-M81I	pMYSH6504 derivative with ATG to ATC mutation at positions 311-313 in <i>virF</i> (ML-397/ML-398)	This study
pF-FS	pMYSH6504 derivative with insertion G after position 208 in <i>virF</i> (ML-331/ML-332)	This study
p _{virB} -lacZ	<i>virB-lacZ</i> transcriptional fusion in pRS415 (bx5/bx6)	This study
pFL-1A	<i>virF-lacZ</i> translational fusion, <i>virF</i> (-289 to +64), from Met1 in frame with the CDS of <i>lacZ</i> (ML-496/ML- FustradFS)	This study
pFL-4A	<i>virF-lacZ</i> translational fusion, <i>virF</i> (-289 to +73) from Met4 in frame with the CDS of <i>lacZ</i> (ML-496/ML- FustradFL)	This study
pFL-M4G	pFL-4A with mutations ATG to GGG at positions 71-73) in <i>virF</i> (ML- ML-296 /ML-297)	This study
pVirF-lacZ	<i>virF(wt)-lacZ</i> translational fusion in pRS414 (pos: -289 to +104) (ML-496/Qh7)	This study
pRS-6504	<i>virF(wt)-lacZ</i> translational fusion in pRS414 (pos: -289 to +405) (ML-496/ML-330)	This study
pRS-M81L	<i>virF(M81L)-lacZ</i> translational fusion in pRS414 (ML-496/ML-330)	This study
pRS-M4G	<i>virF(M4G)-lacZ</i> translational fusion in pRS414 (ML-496/ML-330)	This study
pRS-FS	<i>virF(FS)-lacZ</i> translational fusion in pRS414 (ML-496/ML-330)	This study
pAC-30	pGIP7 derivative carrying <i>virF</i> ₃₀ M81L ORF (ML-333/ML-336)	This study
pAC-21	pGIP7 derivative carrying <i>virF</i> ₂₁ ORF (ML-335/ML-336)	This study
pRS-F(+70)	<i>virF-lacZ</i> transcriptional fusion in pRS415 (pos: +70 to +405) (ML-333/ML-330)	This study
pRS-F(+145)	As pRS-F(+70) but from +145 to +405 (ML-334/ML-330)	This study
pRS-F(+205)	As pRS-F(+70) but from +205 to +405) (ML-1343/ML-330)	This study
pRS-F(+305)	As pRS-F(+70) but from +305 to +405) (ML-1342/ML-330)	This study
pRS-F(+205 -10mut)	pRS-F(+205) derivative with mutations CATTAT to CGTTAT at position +298 to +303 in <i>virF</i> (ML-1344/ML-1345)	This study
pAC-T730-FT	pACYC184 derivative carrying <i>virF</i> sequence (+1) under the control of a T7 promoter (ML-438/ML-566)	This study
pAC-T7-HH	pACYC184 derivative with a T7 promoter and a specific hammerhead ribozyme for <i>virF</i> (ML-U6/ML-U13)	This study
pAC-T7-HH-FT	pACYC184 derivative carrying <i>virF</i> sequence (+309) preceded by a T7 promoter and a specific hammerhead ribozyme sequence (ML-U7/ML-U8)	This study
pGIP7	pACYC184 derivative carrying <i>lacI</i> gene and <i>P_{lac}</i>	(2)
pACYC184	Medium copy number cloning vector	(3)
pRS415	LacZ transcriptional fusion vector	(4)
pRS414	LacZ translational fusion vector	(4)

REFERENCES

1. Sakai T, Sasakawa C, Makino S, Kamata K, Yoshikawa M. 1986. Molecular cloning of a genetic determinant for Congo red binding ability which is essential for the virulence of *Shigella flexneri*. Infect Immun **51**:476–82.
2. Falconi M, Prosseda G, Giangrossi M, Beghetto E, Colonna B. 2001. Involvement of FIS in the H-NS-mediated regulation of *virF* gene of *Shigella* and enteroinvasive *Escherichia coli*. Mol Microbiol **42**:439–52.

3. **Chang AC, Cohen SN.** 1978. Construction and characterization of amplifiable multicopy DNA cloning vehicles derived from the P15A cryptic miniplasmid. *J Bacteriol* **134**:1141–56.
4. **Simons RW, Houtman F, Kleckner N.** 1987. Improved single and multicopy lac-based cloning vectors for protein and operon fusions. *Gene* **53**:85–96.