

*Supporting Information*

# Label-Free Sequence-Specific Visualization of LAMP Amplified *Salmonella* via DNA Machine Produces G-Quadruplex DNAzyme

Huan Zeng <sup>1</sup>, Shuqin Huang <sup>1</sup>, Yunong Chen <sup>1,2</sup>, Minshi Chen <sup>3</sup>, Kaiyu He <sup>2</sup>, Caili Fu <sup>1</sup>, Qiang Wang <sup>2</sup>, Fang Zhang <sup>1,\*</sup>, Liu Wang <sup>2,4,\*</sup> and Xiaohong Xu <sup>2</sup>

<sup>1</sup> College of Biological Science and Engineering, Fuzhou University, Fuzhou 350108, China; 210820068@fzu.edu.cn (H.Z.); 200820041@fzu.edu.cn (S.H.); yunong0420@163.com (Y.C.); caili\_fu@hotmail.com (C.F.)

<sup>2</sup> Institute of Agro-product Safety and Nutrition, Zhejiang Academy of Agricultural Sciences, Hangzhou 310021, China; hekaiyu@zaas.ac.cn (K.H.); wq1357533860@126.com (Q.W.); xuxiahong@zaas.ac.cn (X.X.)

<sup>3</sup> Technology Center of Fuzhou Customs, Fuzhou 350015, China; minshi0271@sina.com

<sup>4</sup> Key Laboratory of Traceability for Agricultural Genetically Modified Organisms, Ministry of Agriculture and Rural Affairs, Hangzhou 310021, China

\* Correspondence: fangzh921@fzu.edu.cn or zhangfang921@gmail.com (F.Z.); wangliually@126.com (L.W.)

## 1. Supplementary Tables and Figures

### 1.1. Supplementary Tables

**Table S1.** Sequences employed in this work.

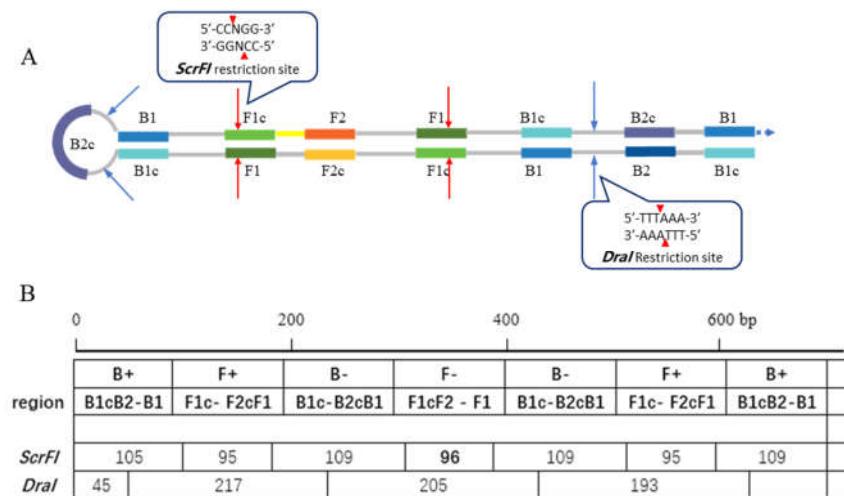
Oligonucleo-tides <sup>a</sup>	Sequence (5'-3') <sup>b,c,d</sup>
FIP	TCCCGGCAGACTTCCCATTGAAATCATGAC-GCAGCTGTTGAA
BIP	TTCCCGCTGCCGGTATTGTTGCTACGTTTGCTTCAC-GGA
F3	GCGATAATATGGGGCGGAAT
B3	CGCCTTGCTGGTTTAGGT
LF	TATTCGGTGGTTTAAGCGTACTC
LB	GCCGTAACAACCAATACAAATGG
Nick-FIP	TCCCGGCAGAG <u>TTCCCATTGAAAGAGTCATCATGAC-GCAGCTGTTGAA</u>
Nick-BIP	TTCCCGCTGCCGGTATTGTT <u>GAGTCGCTACGTTT-GCTTCACCGGA</u>
M-G	<u>TCCCAACCCGCCCTACCCTTTGACTCGGCAGAG-TTCCCATTGAAAT</u>

<sup>a</sup> FIP and BIP are inner primers. F3 and B3 are outer primers. LF and LB are loop primers. Nick-FIP and Nick BIP are FIP and BIP, respectively, incorporated with a nicking endonuclease recognition site. M-G is the molecular machine. <sup>b</sup> Underlined letters indicate the recognition site of Nt.BstNBI. <sup>c</sup> Italic letters in bold indicate the complementary bases of the recognition site of Nt.BstNBI. <sup>d</sup> Underlined letters in bold indicate the complementary sequence of the G-rich sequence.

## 1.2. Supplementary Figures

**F3**  
 781 CGTAAATGGC GATAGCGATA ATATGGGGCG GAATATCATG ACGCAGCTGT TGAACAACCC  
**F2**  
 841 ATTTGTATTG GTTGTTACGG CTATTTGAC CATTCAATG GGAACCTCTGC CGGGATTCCC  
*ScrFI*  
**F1**  
 901 ACTGCCGGTT TTGTTATT TATCGGTGGT TTAAGCGTA CTCTTCTATTT TAAATTCCG  
*DraI*  
**B1c**  
**B2c**  
**B3c**  
 961 TGAAGCAAAA CGTAGCGCCG CCAAACCTAA ACCAGCAAA GGCGAGCAGC CGCTCAGTAT

**Figure S1.** Template sequence for LAMP-Res-Nick amplification of the invA gene of *Salmonella*.



**Figure S2.** The effect of restriction endonuclease, ScrFI, and DraI, on cleaving LAMP products of invA gene of *Salmonella*. (A) Scheme illustrating the restriction sites of ScrFI and DraI on the long stem-loop structure of LAMP products. Red arrows indicate the restriction site of ScrFI. And the blue arrows indicate the restriction site of DraI. (B) The expected LAMP products extended from Nick-FIP. The numbers in the box denote the expected length of the amplification products.