

Supplementary Information to

## **Digital Loop-Mediated Isothermal Amplification on a Commercial Membrane**

Xingyu Lin<sup>1</sup>, Xiao Huang<sup>1</sup>, Katharina Urmann<sup>1</sup>, Xing Xie<sup>1,2</sup>, and Michael R. Hoffmann\*<sup>1</sup>

1. Linde + Robinson Laboratories, California Institute of Technology, Pasadena, California 91125,  
United States

2. School of Civil and Environmental Engineering, Georgia Institute of Technology, Atlanta,  
Georgia 30332, United States

\*Corresponding author, e-mail, [mrh@caltech.edu](mailto:mrh@caltech.edu)

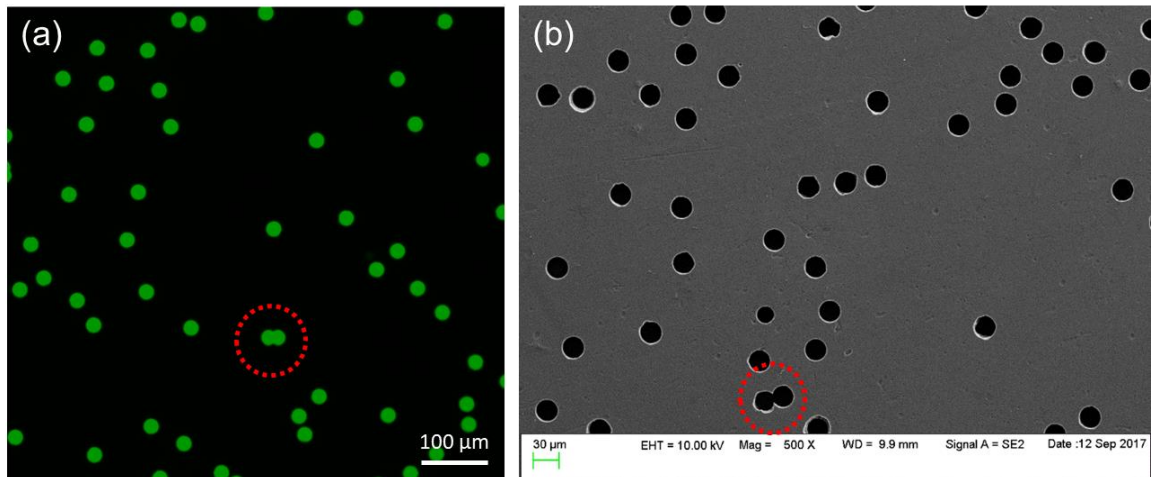
**Table S1.** Primers and probe sequences for mLAMP or mRT-LAMP

	<b>Primer Name</b>	<b>Sequence (5'-3')</b>	<b>Reference</b>
<i>Escherichia coli</i>	F3	GCCATCTCCTGATGACGC	1
	B3	ATTTACCGCAGCCAGACG	
	LF	CTTTGTAACAACCTGTCATCGACA	
	LB	ATCAATCTCGATATCCATGAAGGTG	
	FIP	CATTTTGCAGCTGTACGCTCGCAGCCCATCATGAATG TTGCT	
	BIP	CTGGGGCGAGGTCGTGGTATTCCGACAAACACCACGA ATT	
<i>Enterococcus faecalis</i>	F3	GCCGGAAATCGATGAAGA	2
	B3	TCCAGCAACGTTGATTGT	
	LF	AAATGCTGCGCCAGCTCG	
	LB	TCCAATGTGGAACCTTAAACGTACC	
	FIP	CACTTTTTGTTGTTGGTTTTTCGCTTTATTATCTGCTTGG GGTGC	
	BIP	ATCTGCAGACAAAGTAGTAATTGCTCCAAGCTTTTAA GCGTGTC	
<i>Salmonella Typhi</i>	F3	GACTTGCCTTTAAAAGATACCA	3
	B3	AGAGTGCGTTTGAACACTT	
	LF	TCGGATGGCTTCGTTCTT	
	LB	CAAGGGTTTCAAGACTAAGTGGTTC	
	FIP	AACTTGCTGCTGAAGAGTTGGACCGAATGACTCGACC ATC	
	BIP	CCTGGGGCCAAATGGCATTATGCACTAAGTAAGGCTG G	
MS2	F3	CTTGCGACGATAGACTTATC	4
	B3	TAGATGCCTATGGTTCCG	
	LF	GATTCCGTAGTGTGAGCG	
	LB	GCTAGAGTCCATGATATTCTGG	
	5'FAM -FIP	<b>FAM</b> - ATCGTATCGTCTCGCCATCTA + CCACCAGAGCTATATTCATATC	
	qFIP- 3'IBF Q	GAGACGATACGAT- <b>IBFQ</b>	
	BIP	ACAATGGGAAATGGGTTCACA + GGGTCGCTTTGACTATTG	

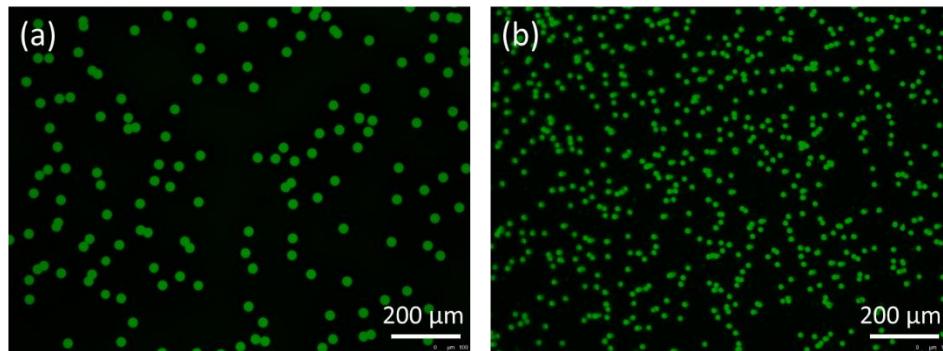
**Table S2.** Primers and probe for qPCR.

	<b>Sequence (5'–3')</b>
Forward primer	5'CGGTGAATACGTTCYCGG3' where Y is either C or T
Reverse primer	5'GGWTACCTTGTTACGACTT3', where W is either A or T
TaqMan probe	FAM-5'CTTGTACACACCGCCCGTC3'

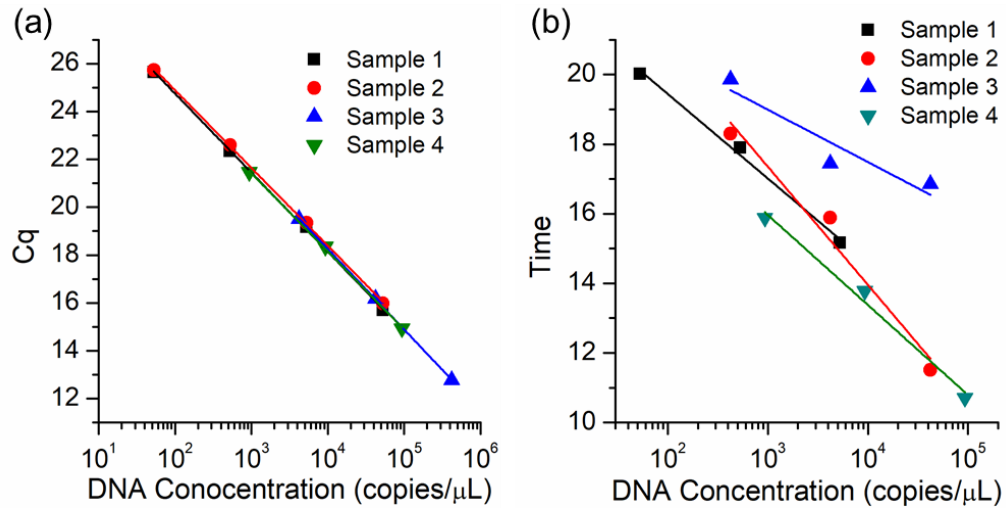
The PCR thermocycling involves 3 minutes of initialization at 95 °C, and 42 cycles of denaturation at 95 °C for 15 seconds, followed by annealing/extension at 55 °C for 30 seconds. The primers and probe are targeting the universal 16s rRNA gene. The sequences are listed above.



**Figure S1.** (a) Fluorescence image of the membrane when pores were filled with fluorescent solution. The red circle indicates overlapped micropores. (b) Top-view SEM image of a membrane. The red circle denotes the overlapped micropores



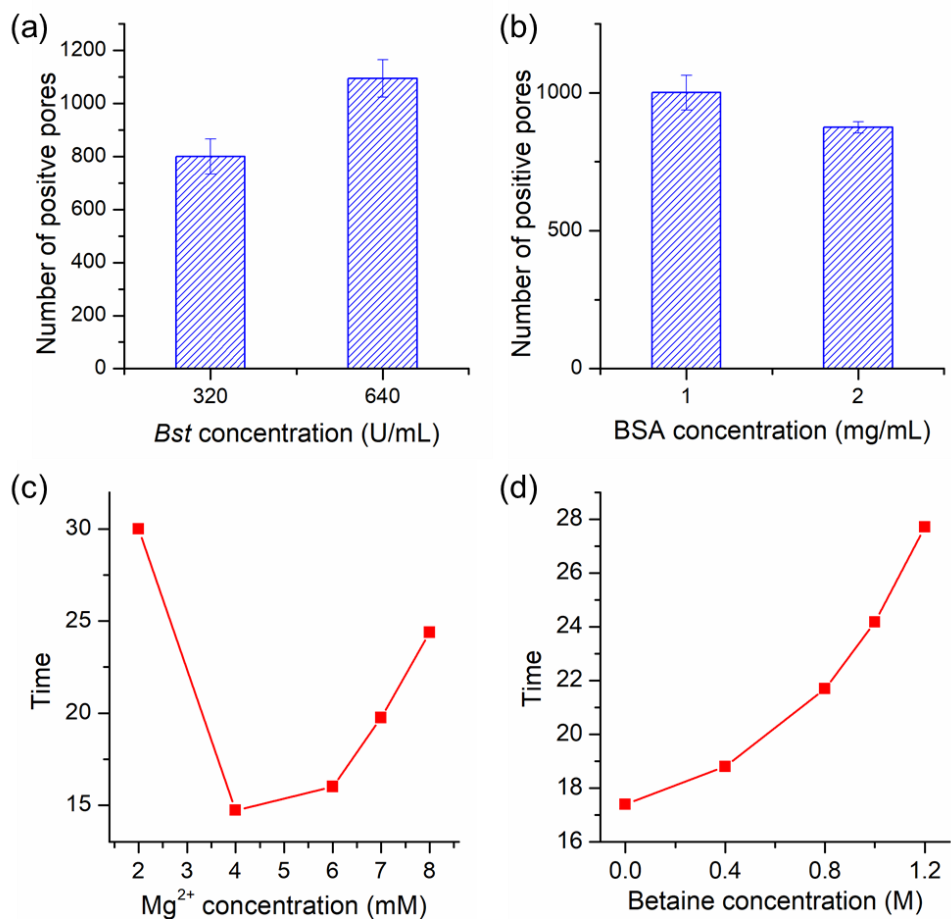
**Figure S2.** Droplet formation on PCTE membranes with different nominal pore sizes. (a) 30  $\mu\text{m}$ . (b) 14  $\mu\text{m}$ .



**Figure S3.** Calibration curves obtained from a series of dilutions of different *E. coli* DNA samples.

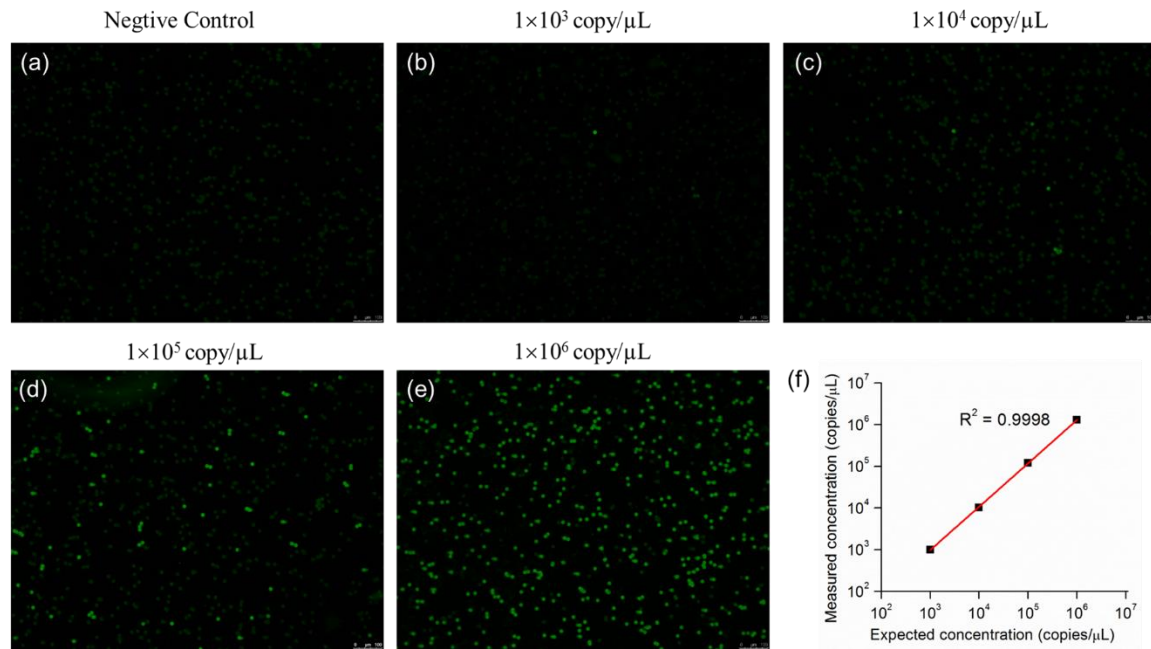
(a) qPCR. (b) qLAMP.

As can be seen in **Figure S3**, the qPCR results based on C<sub>q</sub> show identical calibration curves for different *E. coli* DNA samples. However, the qLAMP results based on T<sub>t</sub> show disordered calibration curves for different samples. Even multiple samples with the same DNA concentration show quite different T<sub>t</sub> values.



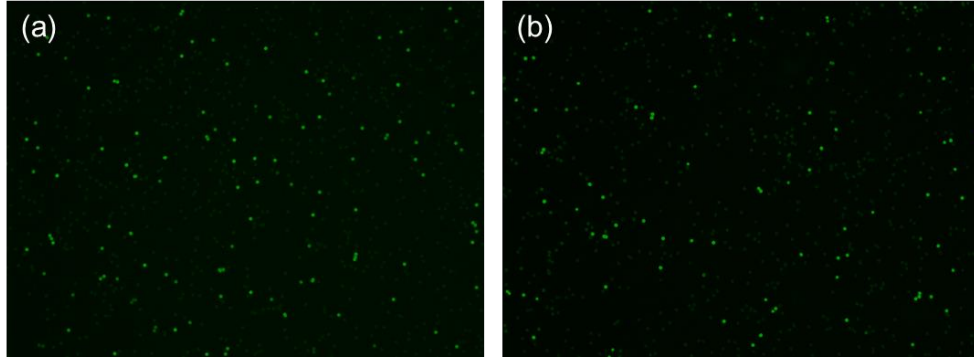
**Figure S4.** Optimization of LAMP reaction. (a,b) Effects of *Bst* (a) and BSA (b) concentrations on the number of positive pores obtained on the membrane. (c,d) Effects of  $Mg^{2+}$  (c) and betaine concentration (d) on Tt value for LAMP performed in the tube. A smaller Tt value means less time is needed for nucleic acid amplification.

According to the optimization results (**Figure S4**), the final 25  $\mu$ L LAMP reactions contained 1  $\times$  isothermal buffer, 6 mM total  $MgSO_4$ , 1.4 mM dNTP, 640 U/mL *Bst* 2.0 WarmStart polymerase, primer mix (1.6  $\mu$ M FIB and BIP, 0.2  $\mu$ M F3 and B3, 0.8  $\mu$ M LF and LB), 1 mg/mL BSA, 50  $\mu$ M calcein, 1 mM  $MnCl_2$  and 2.5  $\mu$ L of template.

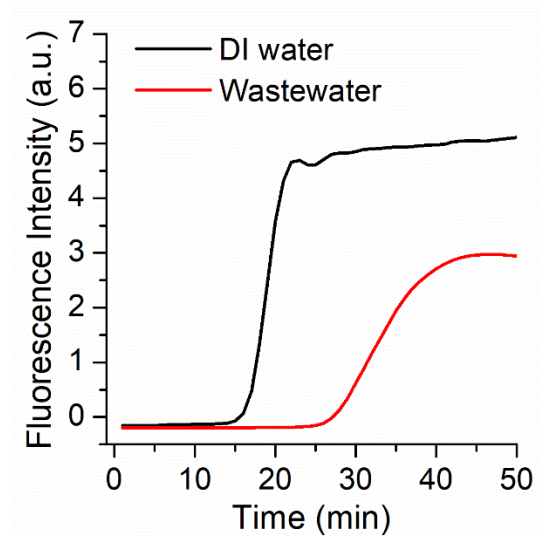


**Figure S5.** (a-e) End-point fluorescence images of membranes (14 μm pore size) after mdLAMP with a series concentration of target *E. coli* DNA. (f) Comparison of measured *E. coli* DNA concentrations to the expected concentrations.





**Figure S6.** mdLAMP analysis of DNA samples extracted from *Enterococcus faecalis* (a) and *Salmonella* Typhi (b).



**Figure S7.** Real-time RT-LAMP results of MS2 quantification in DI water and wastewater.

As shown in **Figure S7**, the qRT-LAMP in the tube was strongly inhibited, resulting in an increased  $T_t$  value.

## Supplementary Movie Legends

**Supplementary Movie S1:** Time lapse movie showing the removal of fluorescent solution on the membrane surface when peeling off the PDMS films. In order to be seen clearly by naked eye, an over-exposure time was used to increase the fluorescence intensity. Therefore, the pores look larger than original.

1. Hill, J.; Beriwal, S.; Chandra, I.; Paul, V. K.; Kapil, A.; Singh, T.; Wadowsky, R. M.; Singh, V.; Goyal, A.; Jahnukainen, T., et al. Loop-Mediated Isothermal Amplification Assay for Rapid Detection of Common Strains of Escherichia Coli. *J. Clin. Microbiol.* **2008**, *46*, 2800-2804.
2. Kato, H.; Yoshida, A.; Ansai, T.; Watari, H.; Notomi, T.; Takehara, T. Loop-Mediated Isothermal Amplification Method for the Rapid Detection of Enterococcus Faecalis in Infected Root Canals. *Oral Microbiol. Immunol.* **2007**, *22*, 131-135.
3. Fan, F.; Yan, M.; Du, P.; Chen, C.; Kan, B. Rapid and Sensitive Salmonella Typhi Detection in Blood and Fecal Samples Using Reverse Transcription Loop-Mediated Isothermal Amplification. *Foodborne Pathog. Dis.* **2015**, *12*, 778-786.
4. Huang, X.; Lin, X.; Urmann, K.; Li, L.; Xie, X.; Jiang, S.; Hoffmann, M. R. A Smartphone Based in-Gel Loop Mediated Isothermal Amplification (gLAMP) System Enables Rapid Coliphage Ms2 Quantification in Environmental Waters. *Environ. Sci. Technol.* **2018**, DOI: 10.1021/acs.est.8b00241.