

Supplementary Document

Point-of-Care Lateral Flow Detection of Viable *Escherichia coli* O157:H7 Using an Improved Propidium Monoazide-Recombinase Polymerase Amplification Method

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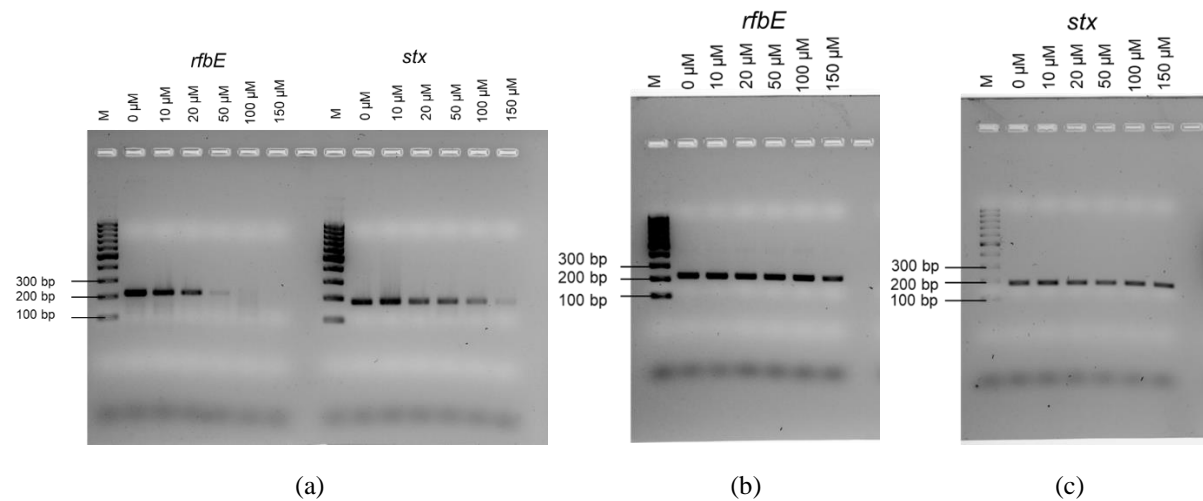


Figure S1: Raw data images for optimisation of the PMAxx-Recombinase Polymerase Assay (RPA) using PMAxx concentrations from 0 to 150 μ M. (a) Lane M: molecular marker, Lane 2 to 7: RPA amplified DNA from dead *E. coli* O157:H7 using *rfbE* F1/R1. Lane 10-15: RPA amplified DNA from dead *E. coli* O157:H7 using *stx* F1/R1. [b&c] PMAxx treated live *E. coli* O157:H7 cells; (b) using *rfbE* F1/R1, (c) using *stx* F1/R1.

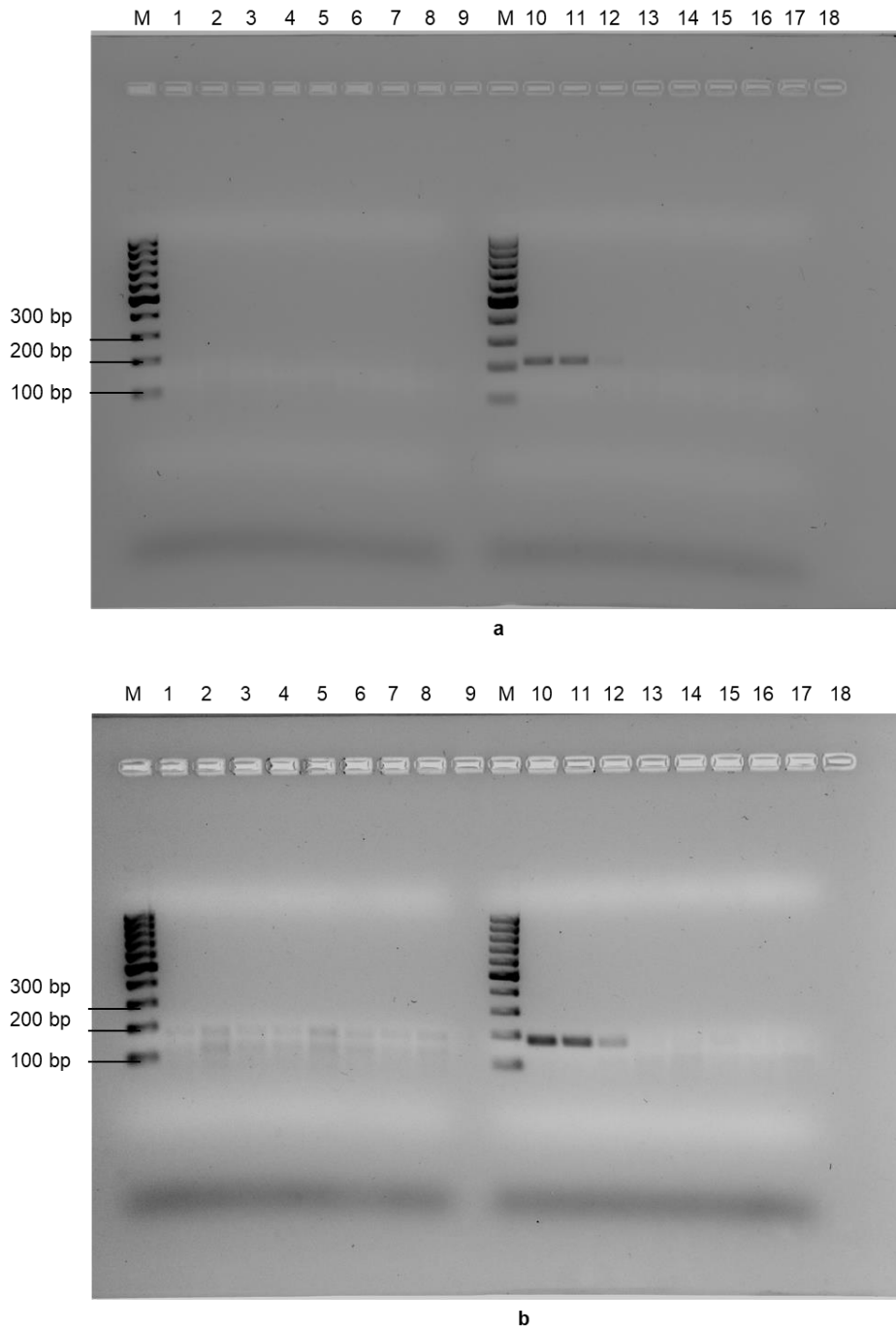
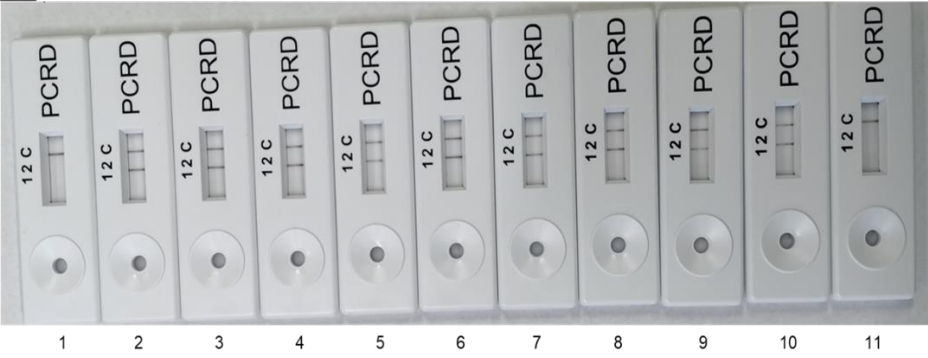
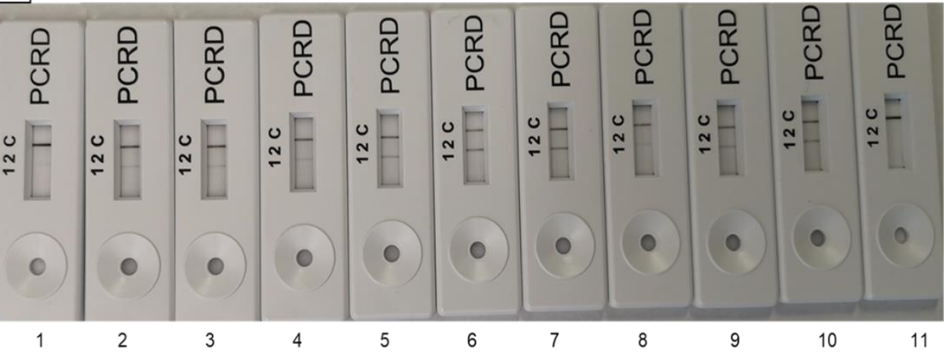


Figure S2: Agarose gel electrophoresis of the PMAxx-RPA assay using different concentrations of *E. coli* O157:H7. (a) targeting *rfbE* gene (b) targeting *stx* gene. Lane M: molecular marker; Lane 1-8: contains dead *E. coli* O157:H7 at concentration 10^8 CFU mL⁻¹, 10^7 CFU mL⁻¹, 10^6 CFU mL⁻¹, 10^5 CFU mL⁻¹, 10^4 CFU mL⁻¹, 10^3 CFU mL⁻¹, 10^2 CFU mL⁻¹, and 10^1 CFU mL⁻¹. Lane 10-18: contains live *E. coli* O157:H7 at concentration 10^8 CFU mL⁻¹, 10^7 CFU mL⁻¹, 10^6 CFU mL⁻¹, 10^5 CFU mL⁻¹, 10^4 CFU mL⁻¹, 10^3 CFU mL⁻¹, 10^2 CFU mL⁻¹, and 10^1 CFU mL⁻¹. PMAxx treatment was given at 100 μ M.

a



b



c



d



Figure S3: Validation of PMAXx- Recombinase Polymerase Assay-Lateral Flow Assay (RPA-LFA) for pure *E. coli* O157:H7 with different concentrations of live and dead cells. (a) and (b) represent the RPA-LFA targeting *rfbE* gene; (c) and (d) represent the RPA-LFA targeting *stx* gene. [a & c]- LFA 1 to 5 PMAXx treated *E. coli* O157:H7 with 0:100, 30:70, 50:50,70:30 ,100:0 live and dead cell ratio. LFA 6 to 10 show control reactions without PMAXx treatment with the same ratio of viable and dead *E. coli* O157:H7. LFA 11 contains NTC. [b & d]- LFA 1 to 5 PMAXx treated *E. coli* O157:H7 with $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$. LFA 6 to 10 show control reactions without PMAXx treatment with the same concentration of live and dead *E. coli* O157:H7 $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$. LFA 11 contains NTC.

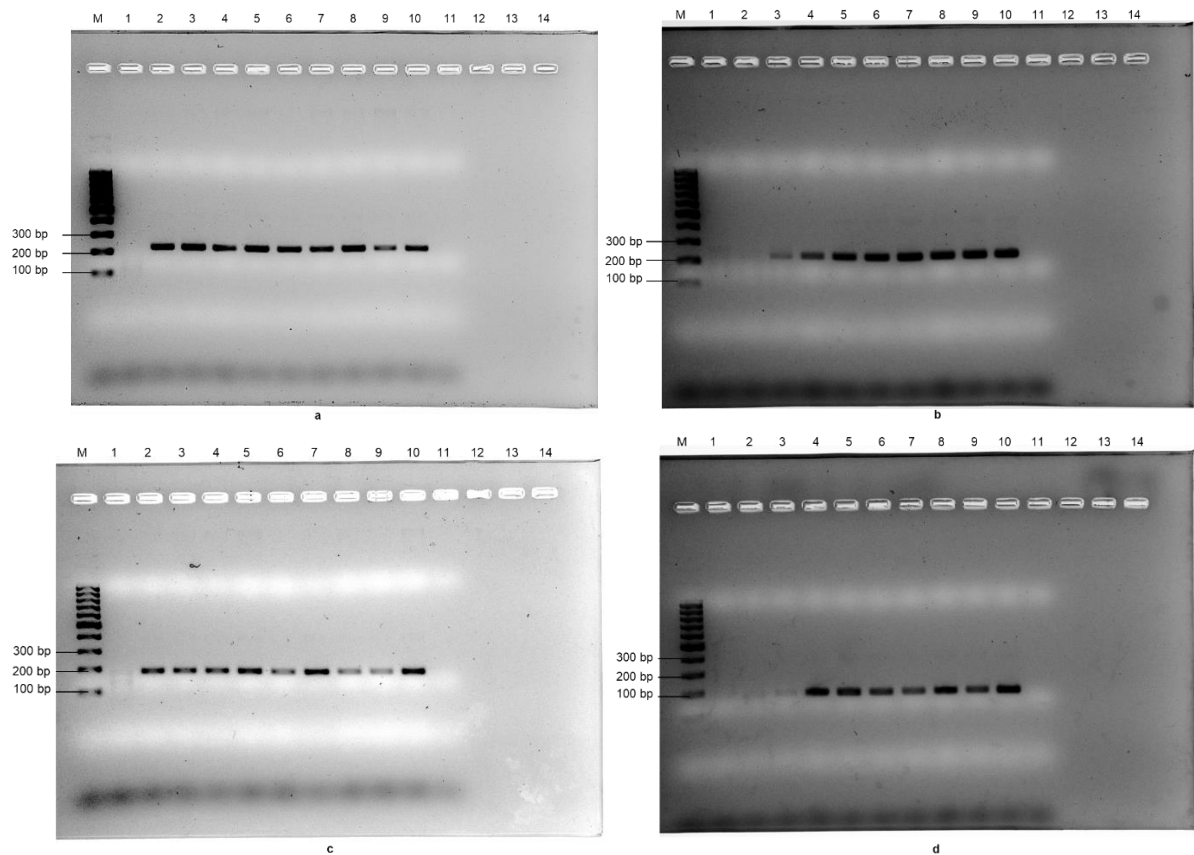


Figure S4: Agarose gel electrophoresis of PMAXx-RPA for pure *E. coli* O157:H7 with different concentrations of viable and dead cells. Image (a) and (b) are representing the RPA assay targeting the *rfbE* gene; image (c) and (d) represent the RPA assay targeting the *stx* gene. [a & c]- lane 1 to 5: PMAXx treated *E. coli* O157:H7 with 0:100, 30:70, 50:50,70:30 ,100:0 viable and dead ratio. Lane 6 to 10 show control reactions without PMAXx treatment with the same ratio of viable and dead *E. coli* O157:H7. Lane 11 contains NTC. Lane M: 100 bp molecular marker. [b & d]- Lane 1 to 5 PMAXx treated *E. coli* O157:H7 with $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$. Lane 6 to 10 show control reactions without PMAXx treatment with the same concentrations of viable and dead *E. coli* O157:H7 $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$. Lane 11 contains NTC.

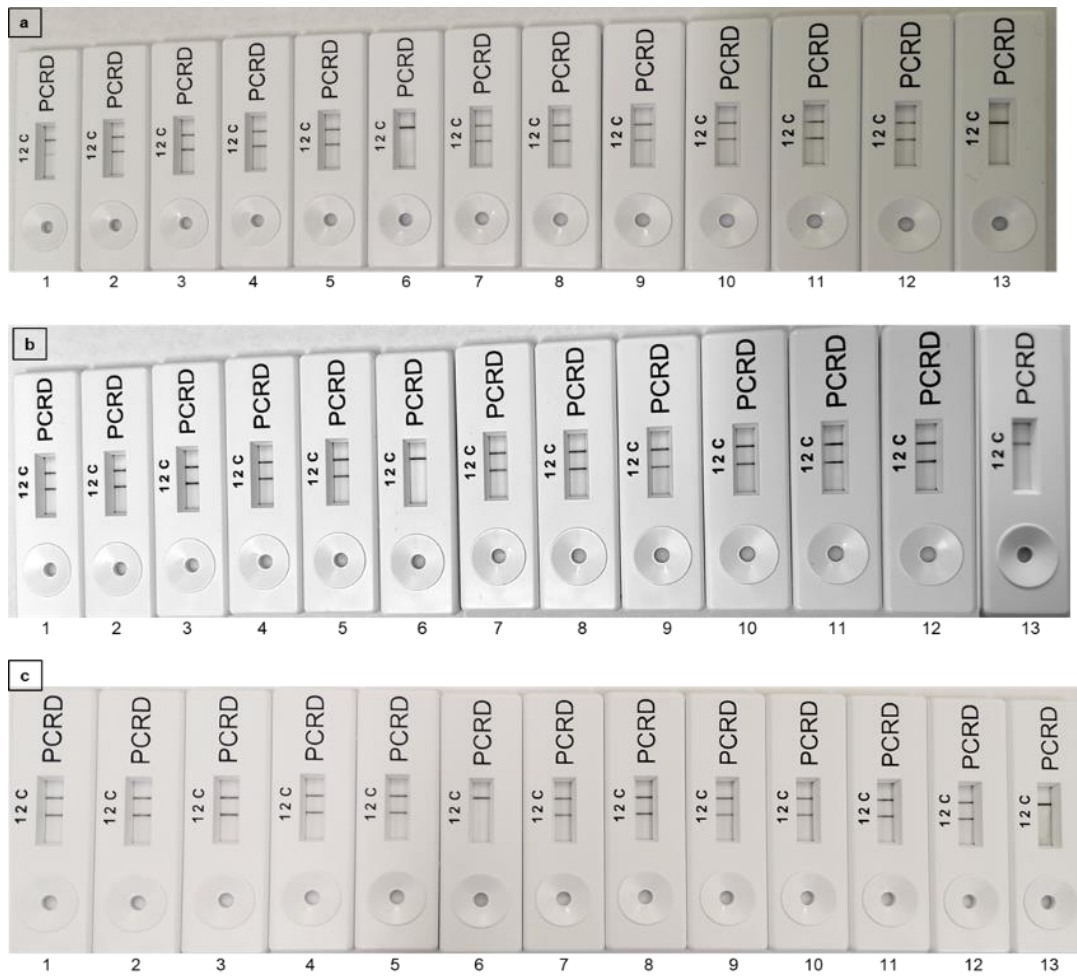


Figure S5: Validation of PMAx- Recombinase Polymerase Assay-Lateral Flow Assay (RPA-LFA) using the *rfbE* gene for commercial beverage samples spiked with different concentrations of live and dead cells, (a) milk samples; (b) water; (c) apple juice. LFA 1 to 6 PMAx treated *E. coli* O157:H7 with $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$, $0:10^8$, live:dead ratio, respectively. Lanes 7 to 12 show control reactions without PMAx treatment with the same concentration of viable and dead *E. coli* O157:H7. Lane 13 shows NTC.

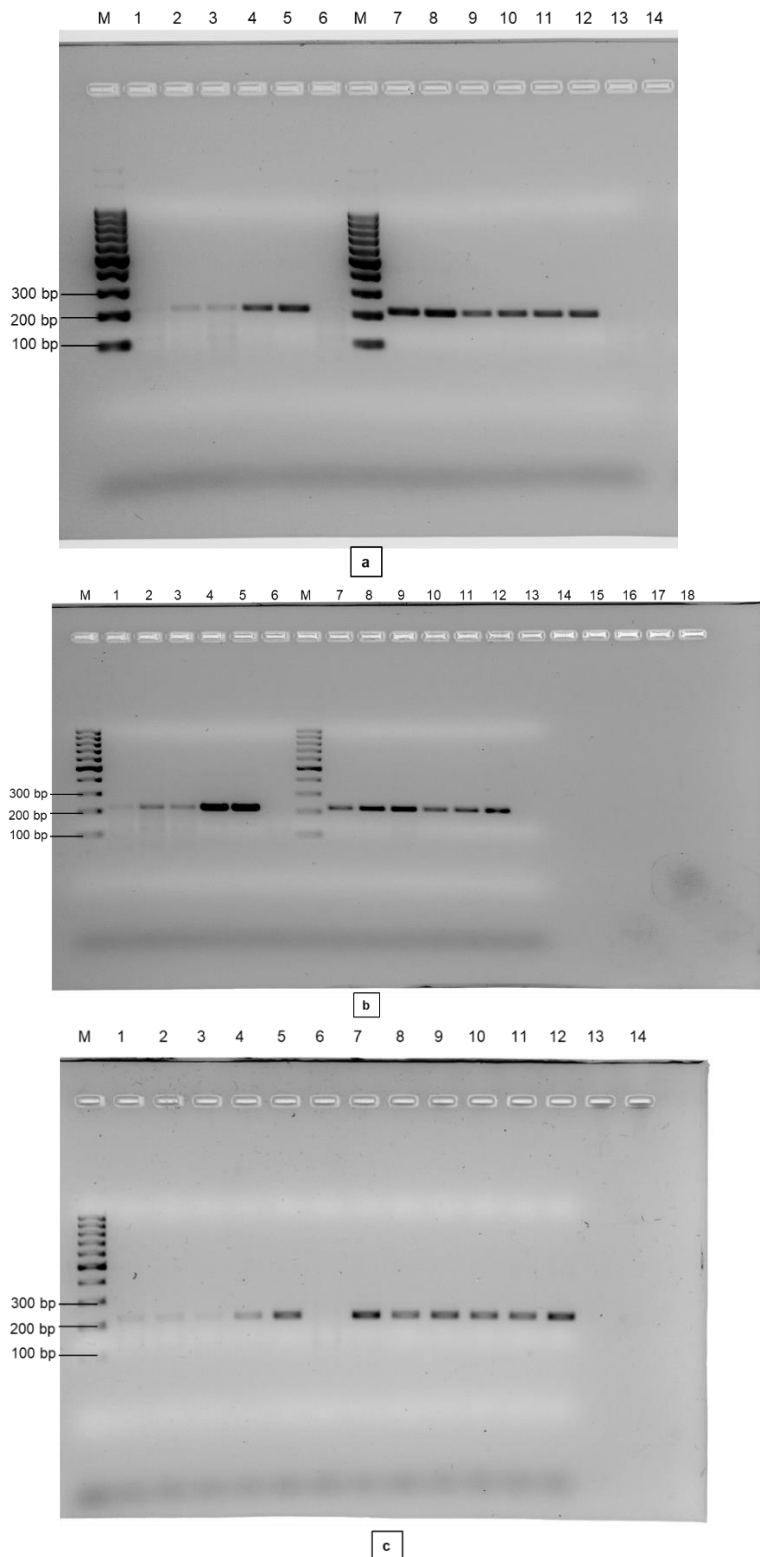


Figure S6: Agarose gel electrophoresis of PMAxx-RPA for commercial beverage samples spiked with varying concentrations of viable and dead cells, (A) milk samples; (B) water; (C) apple juice. Lane M: 100 bp molecular marker, Lane 2 to 6 PMA treated *E. coli* O157:H7 with $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$, $0:10^8$ live and dead. Lanes 7 to 12 show control reactions without PMA treatment with the same concentrations of viable and dead *E. coli* O157:H7 $10^2:10^8$, $10^3:10^8$, $10^4:10^8$, $10^5:10^8$, $10^6:10^8$. Lane 13 contains NTC.

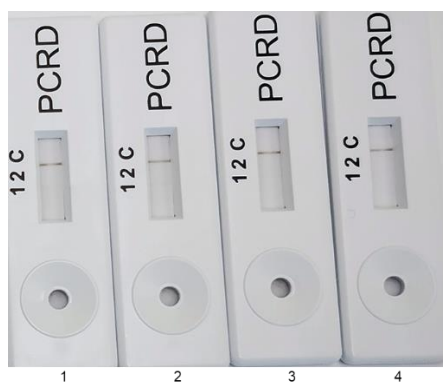


Figure S7: Effect of pH on PMAxx- RPA-LFA for the detection of dead *E. coli* O157:H7. LFA 1: pH 3; LFA 2: pH 5; LFA 3: pH 7; LFA 4: pH 11.

Table S1 Summary of PMAxx- Recombinase Polymerase Assay-Lateral Flow Assay (RPA-LFA) results using different ratios and concentrations of Live:Dead *Escherichia coli* O157:H7 and viable but non-culturable cells (VBNC)

a)

PMAxx-RPA-LFA Live:Dead (CFU/mL)	Gene target <i>rfbE</i>	Gene target <i>stx</i>
(0: 10 ⁸)	✗	✓
(0: 10 ⁷)	✗	✓
(0: 10 ⁶)	✗	✓
(0: 10 ⁵)	✗	✓
(0: 10 ⁴)	✗	✓
(0: 10 ³)	✗	✓
(0: 10 ²)	✗	✓
(0: 10 ¹)	✗	✓
(10 ⁸ :0)	✓	✓
NTC	✗	✗

b)

RPA-LFA Live:Dead ratio (each 10 ⁸ CFU/mL)	PMAxx treatment (<i>rfbE</i> primers)	No treatment (<i>rfbE</i> primers)	PMAxx treatment (<i>stx</i> primers)	No treatment (<i>stx</i> primers)
0:100	✗	✓	✓	✓
30:70	✓	✓	✓	✓
50:50	✓	✓	✓	✓
70:30	✓	✓	✓	✓
100:0	✓	✓	✓	✓
NTC	✗	✗	✗	✗

c)

RPA-LFA Live:Dead (CFU/mL)	PMAxx treatment (<i>rfbE</i> primers)	No treatment (<i>rfbE</i> primers)	PMAxx treatment (<i>stx</i> primers)	No treatment (<i>stx</i> primers)
(10 ² :10 ⁸)	✓	✓	✓	✓
(10 ³ :10 ⁸)	✓	✓	✓	✓
(10 ⁴ :10 ⁸)	✓	✓	✓	✓
(10 ⁵ :10 ⁸)	✓	✓	✓	✓
(10 ⁶ :10 ⁸)	✓	✓	✓	✓
NTC	✗	✗	✗	✗

✗: No coloured band at test line (negative); ✓: coloured band at test line (positive); NTC: non-template control

Table S2 PMA_{xx}-RPA-LFA outcomes in spiked commercial samples

a)

PMA _{xx} -RPA-LFA Live:Dead (CFU/mL)	Milk	Water	Apple juice
(10 ² :10 ⁸)	✓	✓	✓
(10 ³ :10 ⁸)	✓	✓	✓
(10 ⁴ :10 ⁸)	✓	✓	✓
(10 ⁵ :10 ⁸)	✓	✓	✓
(10 ⁶ :10 ⁸)	✓	✓	✓
(0:10 ⁸)	✗	✗	✗
NTC	✗	✗	✗

b)

PMA _{xx} -RPA-LFA VBNC (CFU/mL)	Milk	Water	Apple juice	Pure culture
10 ²	✓	✓	✓	✓
10 ³	✓	✓	✓	✓
10 ⁴	✓	✓	✓	✓
10 ⁵	✓	✓	✓	✓
10 ⁶	✓	✓	✓	✓
NTC	✗	✗	✗	✗

✗: No colored band at test line (negative); ✓: colored band at test line (positive); NTC: non-template control; VBNC: viable but non culturable

Table S3 The unmodified primers for *E. coli* O157:H7 targeting *rfbE* and *stx* gene used for PMA_{xx}-RPA-EF assay

Assay Type	Gene Target	Primer s	Sequence (5'-3')	Primer length (bp)	Amplicon size (bp)	References
RPA- EF	rfbE	Eco rfbE F1	AGCTTTGTTAGCGTTAGGTATATCGGAA GGAGA	33	216	[34]
		Eco rfbE R1	ACATGGATGTCCGTATAAATGGACACAC ATAAT	33		
	stx	Eco stx2 F1	GTGGCCGGGTTTCGTTAATACGGCAACAA ATAC	32	179	
		Eco stx2 R1	TGAAACCAGTGAGTGACGACTGATTTGC ATTC	32		

PMAXx- Propidium monoazide, RPA- recombinase polymerase amplification, EF- electrophoresis.

Table S4 The modified primers and probes for *E. coli* O157:H7 targeting the *rfbE* and *stx* gene used for PMAXx-RPA-LFA.

Assay Type	Gene target	Primers	Sequence (5'-3')	Primer length (bp)	Amplicon size (bp)	References
RPA - LFA	<i>rfbE</i>	Eco rfbE F1	AGCTTTGTTAGCGTTAGGTATATCGGAAG GAGA	33	216	[34]
		Eco rfbE R1	Bt- ACATGGATGTCCGTATAAATGGACACACA TAAT	33		
		Eco rfbE P1 probe	5'-FAM- GTTGATTGAGATAATGAACTTGGCAAAT GT-(THFresidue) TGTTAGTGACATAGAAC- C3spacer-3'	48		
RPA - LFA	<i>Stx</i>	Eco stx2 F1	5'- GTGGCCGGGTTCGTTAATACGGCAACAAA TAC – 3'	32 nt	179	
		Eco stx2 R1	5'- Biotin- TGAAACCAGTGAGTGACGACTGATTTGCA TTC – 3'	32 nt		
		Eco stx2 P1 probe	5'- FAM- AGTGCCCGGTGTGACAACGGTTTCCATGA CAA(THF residue)GGACAGCAGTTATACCA - C3spacer – 3'	49		

PMAXx- Propidium monoazide, RPA- recombinase polymerase amplification, LFA- lateral flow assay

Rani, A., et al., *Evaluation and comparison of recombinase polymerase amplification coupled with lateral-flow bioassay for Escherichia coli O157: H7 detection using different genes*. Scientific reports, 2021. **11**(1): p. 1-12.