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Supplementary information

2 **A Zwitterionic-Enabled One-Pot Electrochemical Strategy for Stable** 3 **End-Point Detection of Isothermal Nucleic Acid Amplification**

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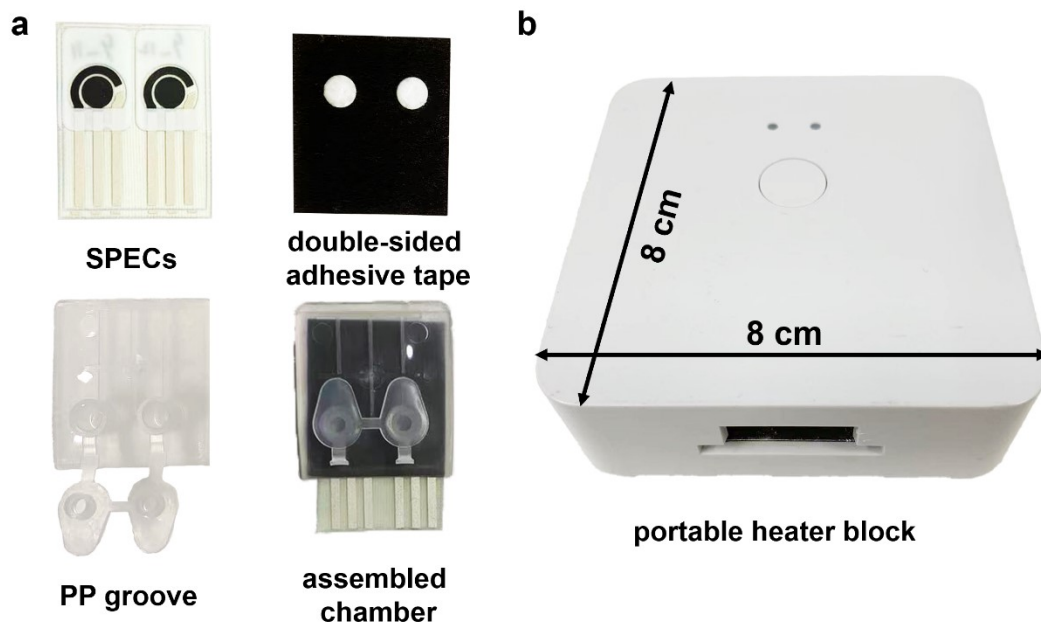
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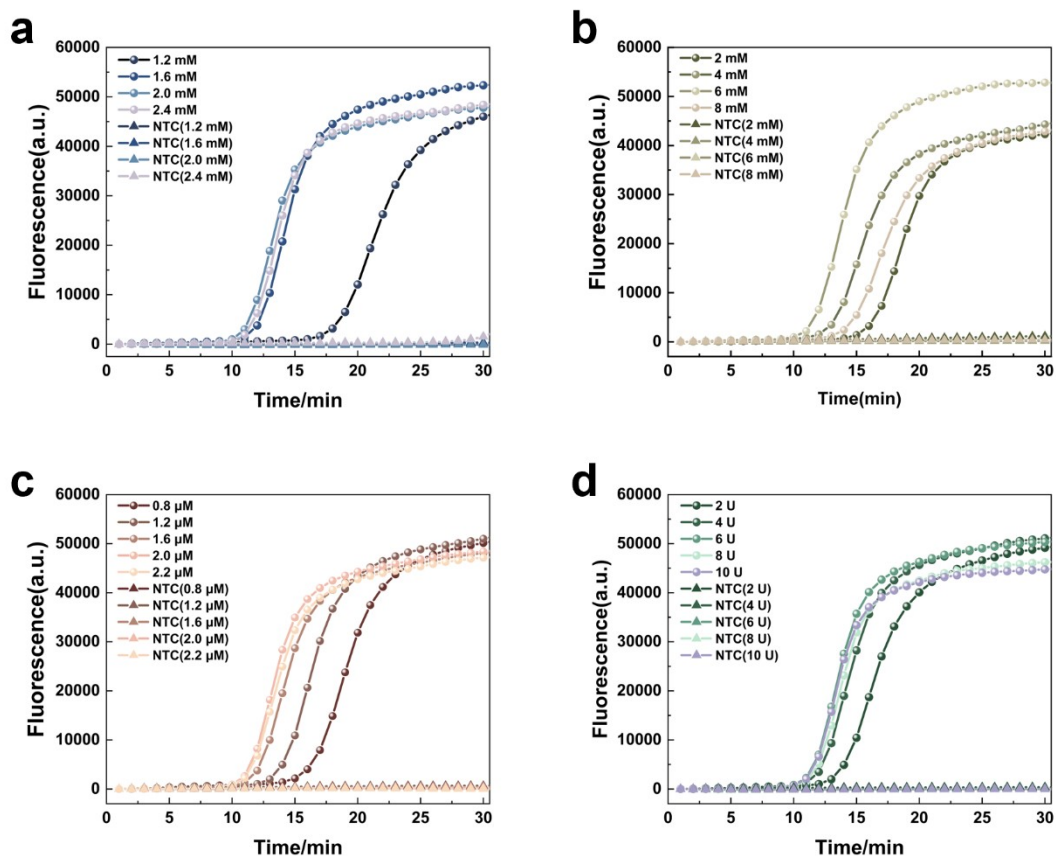
24 **Table S1. Sequences of nucleic acids used in this work**

Name	Sequence (5'-3')
F3	ATGCAAAGAAAATTGAAGTCGA
B3	GCGTTGTCTTCGCTCCAAAT
FIP	CGTTTACCATTTTTCCATCAGCATAGTTTGACAAAGGT CAAAGAACT
BIP	TCAAGGCTTGGCTAAAGTTGCTTATTTTCGCTTGTGCT TCACTT
LF	TACGCTAAGCCACGTCCATA
LB	CCTAACAATACACATGAACAAC
F	AGCGATTGATGGTGATACGGT
R	AGGATGCTTTGTTTCAGGTGT

25 Notes: LAMP primer sequences were adopted from previously published studies¹. PCR primers
 26 were designed and optimized using NUPACK software (<http://www.nupack.org/>) and NCBI
 27 Primer-BLAST (<https://www.ncbi.nlm.nih.gov/tools/primer-blast>).

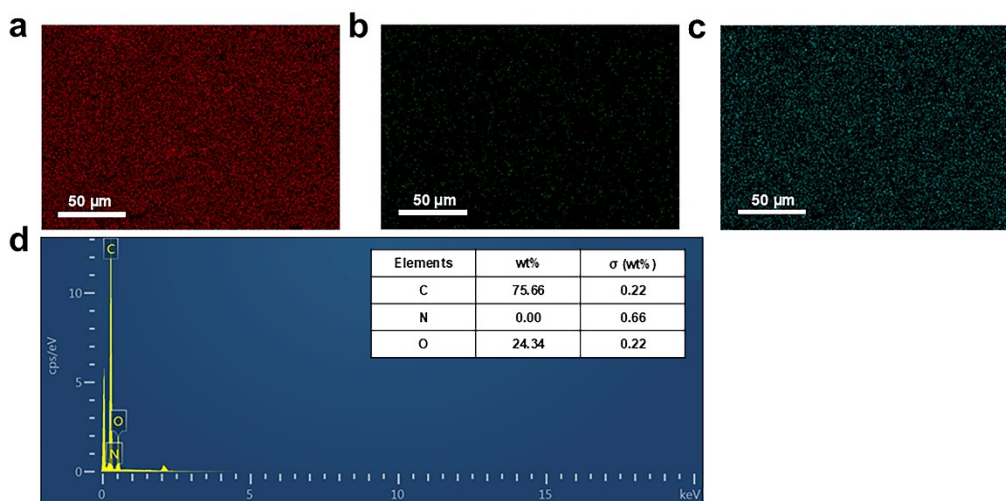


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 29 **Figure S1.** Assembly and configuration of the portable EC-LAMP system. (a) Fabrication of the
 30 reaction chamber by integrating SPCEs, a PP groove, and double-sided adhesive tape. (b) Portable
 31 heater block employed for isothermal incubation during in situ EC-LAMP.



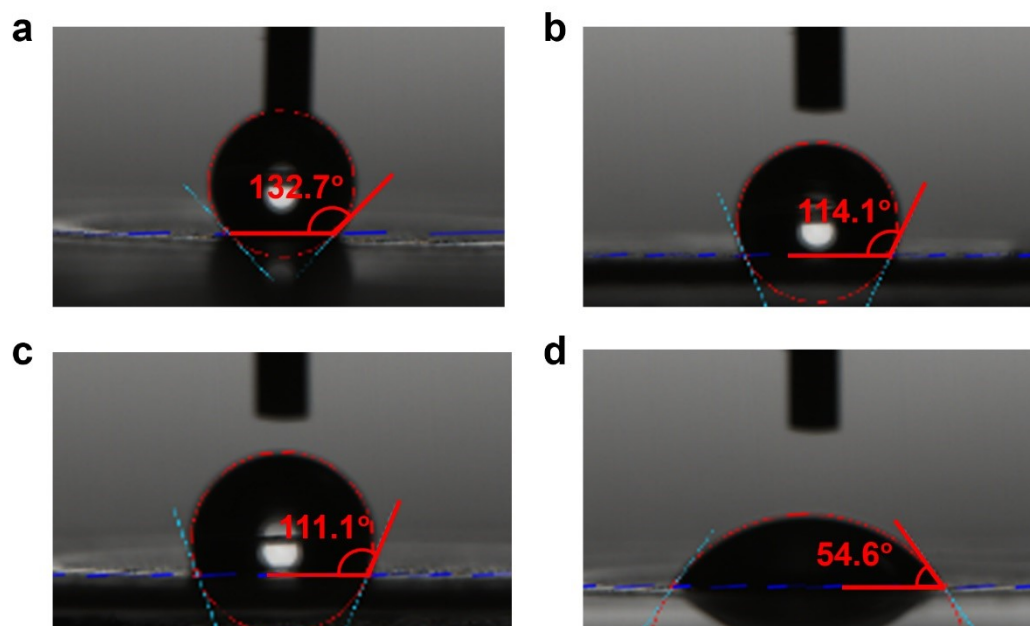
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34 **Figure S2.** Optimization of the reaction conditions for LAMP. (a) different amounts of dNTPs. (b)
 35 different amounts of $MgSO_4$. (c) different amounts of inner primers (FIP/BIP). (d) different
 36 amounts of Bst DNA polymerase.



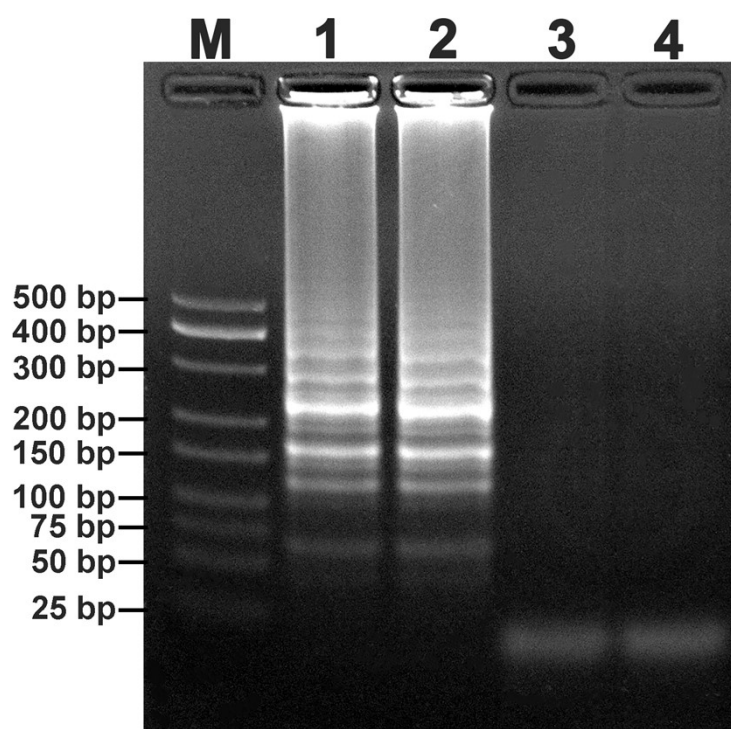
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38 **Figure S3.** (a-c) EDS elemental mapping images of C, N, and O, respectively, of the bare
 39 electrode. (d) EDS spectrum of the selected area, showing the characteristic peaks of C, N, and O.
 40 The inset table lists the corresponding elemental weight percentages (wt%) with standard
 41 deviations (σ).



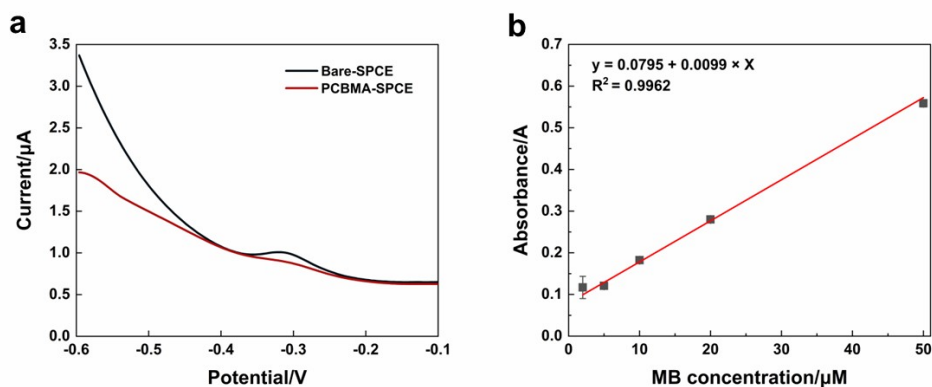
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44 **Figure S4.** Water contact angle images of the electrode surfaces: (a) bare SPCE, (b) H₂SO₄-treated
45 SPCE, (c) EDC/NHS-treated SPCE, and (d) PCBMA- SPCE.



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47 **Figure S5.** Agarose gel electrophoresis image of the LAMP products. Lane 1, Real-time
48 fluorescence LAMP positive; Lane 2, In situ electrochemical LAMP positive; Lane 3, Real-time
49 fluorescence LAMP NTC; Lane 4, In situ electrochemical LAMP NTC. M represents the DNA
50 Marker (25–500 bp). The target concentration was 10⁶ CFU/mL. NTC stands for no target control.



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52 **Figure S6.** Quantitative evaluation of MB adsorption. (a) Electrochemical signals of residual MB
 53 on bare and PCBMA-modified electrodes measured in PBS after removal of MB solution and
 54 triple rinsing. (b) Calibration curve of MB absorbance versus concentration used for quantification.
 55 Error bars represent standard deviations from three independent measurements.

56 To quantitatively evaluate the nonspecific adsorption of methylene blue (MB), the residual
 57 MB concentration in the supernatant after incubation with the electrodes was determined using a
 58 calibration curve (Figure S6b). Based on the Beer–Lambert law, the absorbance values were
 59 converted to concentration using the linear relationship $A = 0.0795 + 0.0099C$. After 30 min
 60 incubation with an initial MB concentration of 20 μM, the remaining concentration (C_e) was
 61 calculated to be 19.90 μM for the PCBMA-modified electrode and 19.46 μM for the bare
 62 electrode. Accordingly, the adsorption amount ($q = C_0 - C_e$) was estimated to be 0.10 μM and
 63 0.54 μM for PCBMA and bare electrodes, respectively. These results indicate that the PCBMA-
 64 modified surface significantly suppresses MB adsorption. The adsorption mitigation efficiency
 65 was calculated to be approximately 81.5%, demonstrating the effective antifouling property of the
 66 zwitterionic PCBMA coating. Although the concentration difference is relatively small, the trend
 67 is consistent with electrochemical measurements.
 68

69 Table S2. Comparison of representative electrochemical LAMP (EC-LAMP)

70 systems reported for nucleic acid detection.

Signal rationale	Electrode / Interface	Antifouling strategy	Time	Sensitivity	Target	References
Potentiometric pH change	Hydrated iridium oxyhydroxide-modified SPCEs	Not found	30 min	$\sim 1.0 \times 10^3$ CFU/mL	<i>Vibrio parahaemolyticus</i>	2
Oxidation of phenol red (pH-dependent)	thick-film carbon electrodes	Not found	122 min	$\sim 2 \times 10^5$ copies/ μ L	<i>Streptococcus pneumoniae</i>	3
pH change via cresol red	3D-printed electrode with pH-responsive readout	UV-laser surface treatment	31 min	~ 11 copies/ μ L	Methicillin-resistant <i>S. aureus</i>	4
Phenol red	SPE graphene-gold-polyaniline (Gr/Au/PANI) electrode	Not found	30 min	20 copies / μ L	<i>SARS-CoV-2</i>	5
pH change	polyaniline (Gr/Au/PANI) electrode	Not found	>30 min	1 copy/ μ L	<i>Helicobacter pylori</i>	6
pH change	polyaniline	Not found	30 min	1 copy/reaction	<i>Mycoplasma pneumoniae</i>	7
Potentiometric pH change	3D printed electrode - modified iridium oxide layer	Not found	>60 min	100 copies/mL	lambda DNA	8
Mediator-	Carbon	Covalent	30 min	$\sim 1 \times 10^3 / 4 \times 10^2$	<i>Staphylococcus</i>	9

displacement	electrode +	probe			copies/reactio	<i>us aureus/</i>	
MB labeled	solid-phase	attachment			n	<i>Treponema</i>	
mediator	universal	reduces				<i>pallidum</i>	
	probe	nonspecific					
	Multiwalled						
Molybdophos	carbon					<i>Nosema</i>	
phate	nanotubes -	Not found	70 min		~17 fg/μL	<i>bombycis</i>	10
precipitate	modified						
	GCE						
molybdophos					~1.0 × 10 ³		
phate	SPCEs	Not found	45 min		CFU/mL 5.29	SARS-CoV-2	11
precipitate					copies/μL		
Digoxigenin-	Carbon					PCA3	
dUTP	electrode	Not found	110 min		Not found	lncRNA	12
	chips						
	a disposable				3 × 10 ²		
Hoechst	electrochemic	Not found	60 min		copies/reactio	GMO	13
33258	al printed				n		
	(DEP) chip						
						plasmid DNA	
RuHex	gold	Not found	60 min		10 ¹ copies	(parvovirus	14
	electrode					VP2 gene)	
Malachite	Thick-film				~22	SARS-CoV-2	15
green	screen printed	Not found	30 min		copies/μL		
	electrodes						
Methylene	AuNPs/SH-				~17.7	<i>Salmonella</i>	
blue	DNA probe/	Pyrene	40 min		CFU/mL	<i>enteritidis</i>	16
Methylene					~2.5 × 10 ⁻⁶	SARS-CoV-2	17
blue	SPCEs	Not found	30 min		ng/μL		
Methylene	PCBMA-					<i>Staphylococc</i>	
blue	modified	PCBMA	30 min		12 CFU/mL	<i>us aureus</i>	This work
	SPCEs						

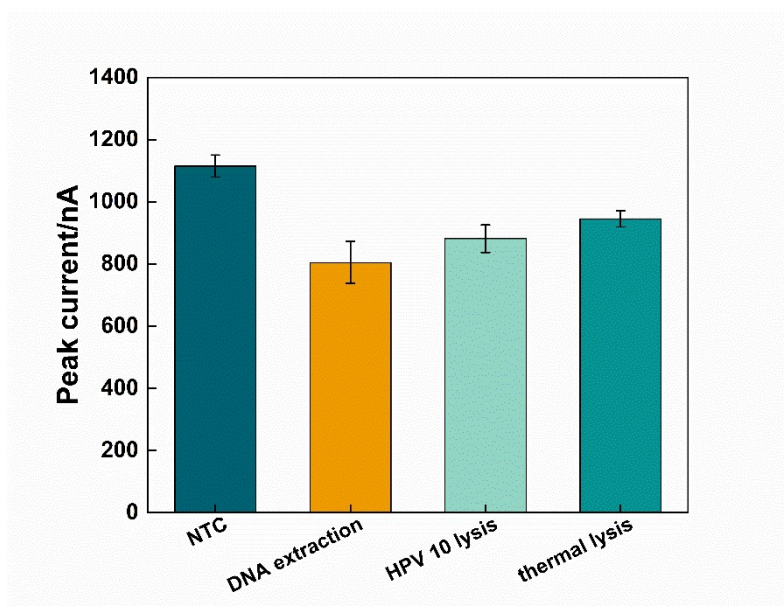
73 **Table S3. Determination of *S. aureus* in milk samples with the proposed**
 74 **biosensor.**

Spiked concentration (CFU mL ⁻¹)	Found concentration (CFU mL ⁻¹) #	Recovery (%)	RSD (%)
1.0 × 10 ²	9.04 × 10 ¹	90.4	0.66
1.0 × 10 ⁴	9.18 × 10 ³	91.8	1.44
1.0 × 10 ⁶	9.69 × 10 ⁵	96.9	2.13

75 #Average of three measurements

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79 **Figure S7.** Electrochemical responses obtained from spiked milk samples (10³ CFU mL⁻¹)
 80 subjected to different pretreatment methods. NTC: no-template control; DNA extraction:
 81 commercial kit-based purification; nucleic acid releaser (95°C, 5 min); thermal lysis: direct
 82 heating (95°C, 5 min).

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