



Loop-Mediated Isothermal Amplification (LAMP) as a Rapid, Affordable and Effective Tool to Involve Students in Undergraduate Research

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Undergraduate research (UR) is a high-impact practice (HIP) to engage undergraduate student in science, technology, engineering and mathematics (STEM), especially from underrepresented groups. UR experiences (UREs) can be integrated into the classroom, making authentic research experiences inclusive and available to all students. However, developing UR pedagogy can be challenging for faculty in resource-limited labs, such as community colleges and small liberal arts colleges. Often molecular biology research methods are expensive, time-consuming and need equipment not readily available or affordable in small schools. Polymerase chain reaction (PCR) is one of the most commonly used techniques in research labs and many UREs. We have investigated loop-mediated isothermal amplification (LAMP) as an inexpensive, accessible alternative to PCR for DNA amplification enabling the identification of microorganisms in the context of UREs. LAMP does not require expensive instrumentation or reagents and uses equipment commonly found in teaching labs. By performing the technique, students learn several key scientific skills that will be useful in their undergraduate or graduate STEM careers. We designed guided independent research experiences for several undergraduates that included the use of LAMP. Students successfully applied the technique to culture samples of common environmental bacteria, including Escherichia coli, Salmonella spp., Staphylococcus aureus, and Enterococcus, and were in addition, able to detect both Salmonella and Enterococcus in directly sampled environmental waters. To highlight the accessibility and affordability of this URE, a simple boiling method was used for DNA preparation from environmental samples. Student response

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data show positive attitudes toward UR when LAMP is utilized as a research tool to tackle relevant biological questions. The feasibility of using simplified LAMP in UREs points to a potential, more expanded application to public engagement with science and broader and more inclusive interactions with the research community.

Keywords: LAMP, research tool, undergraduate research (UR), HIP's, stem education

INTRODUCTION

The Association of American Colleges and Universities has recognized several instructional modalities as high-impact practices (HIPs) that increase student retention rates and student engagement (Kuh, 2008). Among these practices, undergraduate research (UR) has been found to be a particularly effective pedagogy for the engagement and retention of undergraduate students (Kuh, 2008). UR can provide students with not only scientific skills but also increased self-confidence, improved oral and written presentation skills and enhanced critical thinking (Kuh and O'Donnell, 2013).

Two main challenges are faced when creating UR experiences (UREs). First, it may be difficult to design projects that are realistically feasible and that also engage students in relevant authentic queries. Second, many procedures require reagents, equipment and/or central technical support that are beyond the means of many undergraduate institutions. One category of studies that can address the first issue consists of projects that focus on environmental sampling, such as the identification of pathogens from environmental samples like soil and water. The current COVID-19 pandemic has heightened student awareness of the value of such studies. However, environmental pathogen identification typically involves the use of polymerase chain reaction (PCR) for DNA amplification, which is time consuming, expensive and can be difficult for students to perform accurately. At Queensborough Community College (QCC) we have developed UREs in which loopmediated isothermal amplification (LAMP) can be used as a simpler, sensitive and far less expensive alternative to PCR for the identification of microbes in samples obtained from locations on campus as well as from the surrounding urban environment. While comparable in its value as a teaching tool to PCR, LAMP is over tenfold less expensive, as well as more sensitive and more robust in handling complex biological samples (Kaneko et al., 2007; Mori and Notomi, 2009; Law et al., 2014; Warghane et al., 2017). The simplicity of colorimetric readouts, coupled with the simplicity of the amplification procedure itself, makes it easy to incorporate LAMP in a classroom context, so that the method can be used to support course-based UREs (CUREs) as well as independent UR projects.

The standard LAMP assay does not require equipment beyond those that are available in standard biology laboratories (see **Supplementary Figure S1**). The simplicity of the setup derives from the dependence of LAMP on the inherent strand-displacing activity of the Bst polymerase, avoiding the need for the repeated high-temperature intervals employed in PCR, so that the entire reaction can be performed at a single temperature (60–68°C), using a non-denatured doublestranded DNA template (see **Supplementary Figure S2**). However, sample preparation methods prior to the assay may still consume classroom time and require specialized equipment and/or expensive supplies. By using the simple boiling method of sample preparation, LAMP can be made accessible not only to on-campus undergraduates but also to the at-home student in the context of distance learning, now emerging as a dominant mode of instruction during the COVID-19 pandemic. Even beyond the at-home student, we suggest that simplified LAMP protocols that avoid or minimize sample preparation may be uniquely positioned to promote a broader scientific engagement with lay communities and foster enhanced interactions between the research community and the public.

We describe here some initial illustrative data generated by several of our undergraduates with whom we designed guided UREs. These students were able to use LAMP to detect several bacterial microorganisms commonly found in the environment: *Escherichia coli (E. coli), Salmonella* spp. (*S.* spp.), *Staphylococcus aureus (S. aureus)*, methicillin-resistant *Staphylococcus aureus* (MRSA), and *Enterococcus* (Hill et al., 2008; Tang et al., 2012; Wang et al., 2015; Lin et al., 2017; Hu et al., 2018).

MATERIALS AND METHODS

Setting the URE

The students who participated part in the URE were either Biotechnology or Chemistry majors in an urban community college who had enrolled in a research laboratory internship course. Students had taken two semesters of introductory biology as well as biotechnology either prior to or concurrent with the URE.

Bacterial Strains and Growth Media

Escherichia coli, Salmonella spp., *Staphylococcus aureus, Enterococcus faecalis*, and methicillin-resistant *Staphylococcus aureus* (MRSA) were grown in Brain Heart Infusion Broth or Tryptic Soy Broth.

Sample Preparation

Soil samples (approximately 5 ml) were suspended in 50 ml water in conical tubes and allowed to settle prior to removal of 10 ml supernatant for DNA extraction by the boiling method (De Medici et al., 2003). DNA extraction from standing water samples (10 ml) was carried out by the boiling method.

	Target gene	Incubation temperature	References
E. coli	malB	66°C	Hill et al. (2008)
Salmonella spp.	fimY	65°C	Tang et al. (2012)
Salmonella enteritidis	safA	65°C	Azinheiro et al. (2018)
S. aureus	nucAfemA (MRSA)	60°C60°C	Wang et al. (2015)Lin et al. (2017)Chen et al. (2017)
Enterococcus spp.	23S rRNA	64°C	Martzy et al. (2017)

For *Enterococcus* tests, samples (10 ml) were collected from the East River (40.80557°N, 73.79661°W) and extracted by the boiling method.

DNA preparation was either by column-based purification (Quick-DNA Microprep Kit, Zymo Research, Irvine, CA, United States) or by the boiling method (De Medici et al., 2003). For the boiling method, all samples were first spun and resuspended at $1/10^{th}$ original volume.

LAMP Assay

The target genes and protocols employed in the study are summarized in **Table 1**. Note that *fimY* (encodes for the Fimbriae Y protein) and *nucA* (encodes for the nuclease) are conserved targets for all serotypes of *Salmonella* and *S. aureus*, respectively, while *safA* (encodes for the major subunit of *S. enterica* atypical fimbriae) is specific for *Salmonella enteritidis. malB* (encodes for maltose operon protein B) is specific to *E. coli* and *femA* (encodes for methicillin-resistant *S. aureus* (MRSA).

All LAMP incubations were for 1 hour. Reaction products were visualized by ethidium bromide staining of agarose gels. The amplified products from the LAMP reaction are not singlesize amplicons but rather exhibit a range of different product sizes (**Supplementary Figure S2**, Part 5b). Thus, a positive LAMP reaction appears as a smear or a ladder of amplified products on an agarose gel, rather than a single band as seen with PCR (Hill et al., 2008; Tang et al., 2012; Wang et al., 2015; Lin et al., 2017; Hu et al., 2018). For detection of *Enterococcus* spp. in environmental samples, the LAMP amplification of the 23S rRNA target was performed with the addition of SYBR Green I, and the reaction was performed for 60 min in a qPCR machine (Quantstudio 6, Applied Biosystems in order to quantitate fluorescence each minute as a measure of product yield).

Detection of Enterococcus by Enterolert

The Enterolert test (IDEXX Laboratories, Westbrook, Maine) is an enzymatic reaction based on the ability of *Enterococcus* beta-glucosidase to cleave 4-methyl-umbelliferyl b-D-glucoside, yielding a fluorescent product. Briefly, water samples for enumeration of *Enteroccoci* via IDEXX Enterolert Media were collected from the East River (40.80557°N, 73.79661°W) in 1 L acid washed high-density polyethylene bottles. Samples were processed the same day per the manufacturer's protocol. After incubation at 41°C for 24 h, samples were read using a 6-watt, 365 nm UV light box, and enumerated using the IDEXX MPN table.

Laboratory Safety

The URE in which students participated includes education in the biology and clinical impact of the pathogens studied. Standard biosafety practices for microbiology laboratory work were enforced, including adherence to Biosafety Level 2 (BSL2) procedures. Aseptic technique with a Bunsen burner was used for bacterial work. No enrichments of environmental samples were performed, although students were made aware that these samples can contain disease-causing organisms.

RESULTS

Amplification From DNA of Known Strains

Optimization

As a first step toward the incorporation of LAMP in an URE, students performed amplifications using column-purified DNA prepared from cultures of *E. coli*, *S. aureus*, *Salmonella* spp., and *Enterococcus* spp. As shown in **Figures 1–4**, successful amplification was obtained with all four strains. The *E. coli* and *S. aureus* amplifications were used as a basis for optimization studies, and students were able to demonstrate a range of product



FIGURE 1 | LAMP amplification of *E. coli.* (A) Temperature optimization of *E. coli* LAMP. Lane 1, lambda *Hin*dIII marker. Lanes 2, 4, and 6 were generated with chromosomal DNA and Lanes 3, 5, and 7 with water as negative control. Lanes 2 and 3 were incubated at 70°C, Lanes 4 and 5 at 66°C and Lanes 6 and 7 at 62°C. (B) Specificity of *E. coli* LAMP. Lane 1; *E. coli*, Lane 2; *S. typhimurium*, Lane 3; *Staphylococcus aureus*, Lane 4; water, Lane 5; 1 Kb marker. All reactions were carried out at 66°C.



yields by temperature variation, with optimal results within the 60–66°C range as expected (**Figures 1**, **2**). LAMP affords multiple additional parameters for student design of optimization studies, including inner/outer primer ratio, magnesium concentration and incubation time, which will be incorporated in future development of the URE.

Specificity

Students were able to confirm the specificity of LAMP for *E. coli* (Figure 1), *Salmonella* spp. (Figure 3), and *Enterococcus* (Figure 4), as well as the conservation of the *fimY* target in multiple *Salmonella* strains (Figure 3). Students were also able to observe non-specific elements, such as primer bands and chromosomal DNA (Figure 1B, lane 2). An ambiguous result was obtained with *Streptococcus* (*Enterococcus*) (Figure 3A, lane 5), likely reflecting contamination and/or excess chromosomal DNA. The exquisite sensitivity of LAMP enhances the opportunity for students to observe such false positives, providing opportunity for critical thinking and troubleshooting.





DNA-Preparation Method

To assess the dispensability of column-based DNA preparation in the pedagogical use of LAMP, students were given a blind series of sixteen distinct *S. aureus* strains, of which six were MRSA, and used both column and boiling methods for each strain, prior to performing *mecA* LAMP to identify MRSA strains. Similar performance was observed with the two DNA preparation methods: in each case, two MRSA



control). (C) LAMP amplification of S. enteritidis (safA target). Lane 1, Lambda HindIII marker. Lane 2, S. enteritidis. Lane 3, water.



strains were correctly identified (one of which was identified by both methods) and in each case there were four false negatives (data not shown). Overall, these data are supportive of the utilization of LAMP in classroom-based and distance learning settings.

Amplification From Environmental Samples

Our students used the LAMP assays they had established to assess environmental samples for either Salmonella spp. or Enterococcus spp. Enrichments were not performed, since the goal was to determine whether LAMP assay was adequate to allow students to detect these microorganisms with minimal processing. Salmonella spp. was detected in a water sample from a kettle pond in an urban park (Alley Pond, Bayside, NY, United States), as well as in standing water and soil samples within the college campus (Figure 5). Enterococcus was assessed by LAMP in a series of samples taken over several months from three locations in an urban waterway (East River in New York City). Enterococcus levels in these samples were also quantitatively assessed by defined substrate technology (Enterolert; IDEXX). To allow the students to more objectively relate the Enterolert data to their qualitative LAMP findings, the LAMP results were made semi-quantitative by determining the reaction time (T_t) required to achieve a threshold level of SYBR Green fluorescence. The T_t values were used to divide the samples into a LAMP-High group ($T_t < 22$) and a LAMP-Low group ($T_t > 30$). The samples were also divided on the basis of the Enterolert data into groups that were either high or low for most probable number (MPN, comparable to CFU). The MPN-High group had MPN values of 110-3,578) while the MPN-Low group had MPN <10-30. It was observed that seven of eight

Sample	Date	MPN	LAMP		Sample	MPN	LAMP		
IT1	5/23/2019	3578	High		IT1	3578	High	٦	:
HC1	5/23/2019	<10	High		HC2	2282	High	1	
MAR1	5/23/2019	354	High		HC4	1565	High		
IT2	5/30/2019	1515	High		IT2	1515	High		
HC2	5/30/2019	2282	High	MPN-High	< IT4	402	Low		
MAR2	5/30/2019	145	High		MAR1	354	High		
IT3	6/6/2019	20	Low		MAR2	145	High		
HC3	6/6/2019	10	High		MAR5	110	High		
MAR3	6/6/2019	10	Low			34		J	
IT4	6/13/2019	402	Low		HC5	30	Low		- p = 0.02
HC4	6/13/2019	1565	High		IT3	20	Low		
MAR4	6/13/2019	10	Low		IT5	20	High		
IT5	6/20/2019	20	High	<i>4</i>	HC6	20	Low		
HC5	6/20/2019	30	Low		HC3	10	High		
MAR5	6/20/2019	110	High	MPN-Low	MAR3	10	Low		
IT6	6/27/2019	<10	Low		MAR4	10	Low]	
HC6	6/27/2019	20	Low		MAR6	10	Low		
MAR6	6/27/2019	10	Low		HC1	<10	High] [
1	1		1		IT6	<10	Low]	
								- J	

	Power supply (\$600)	Electrophoresis tank (\$400)	UV transilluminator (\$1,000–5,000)	Thermocycler (\$3,000)	Water bath (\$500)	Total cost
	(\$000)	unit (\$ 100)	(\$1,000 0,000)	(\$0,000)	(\$555)	
PCR	\checkmark	\checkmark	\checkmark	\checkmark	—	\$5,500
LAMP	\checkmark	\checkmark	\checkmark	_	\checkmark	\$2,500
Simple LAMP*	_	_	_	_	\checkmark	\$500
At-home LAMP*	_	_	_	_	_	\$0

TABLE 2 | Materials required to set up LAMP and PCR.

*LAMP using colorimetric readout and samples prepared by boiling.

MPN-High samples were LAMP-High, compared to three of ten samples in the MPN-Low group. This difference was significant at p = 0.02 by Fisher's exact text (**Figure 6**). These results show that students are able to apply LAMP semi-quantitatively to assess environmental levels of bacterial pathogens as part of an URE.

DISCUSSION

We sought a new approach to developing UR projects to identify common microorganisms that can be detected in environmental samples. Standard methods for the detection of these microorganisms involve culturing and isolating the bacteria, which is a time-consuming and laborious process. Amplification of genomic DNA by PCR is considerably faster but requires an expensive thermocycler and somewhat expensive reagents. We propose LAMP as a promising alternative method to detect microorganisms, as it is rapid, inexpensive with respect to both equipment and reagents, and does not require much effort to set up (**Tables 2, 3** and **Supplementary Figure S1**).

We demonstrate here that, even in the undergraduate setting, the LAMP technique can be effectively used to replace PCR to help identify bacterial microorganisms. Our data demonstrate that students with modest training in molecular biology can be involved in authentic research aimed at detecting microbes in unknown environmental samples. We have trained eleven students over the course of three semesters to successfully use this technique to detect environmental microbes. We used LAMP to amplify four common and clinically relevant environmental bacteria: *E. coli*, *S. enteritidis, Enterococcus*, and *S. aureus*. All of the data presented here were generated in student-performed studies conducted within a semester time frame, as part of an independent study research course. The students not only acquired basic laboratory proficiency but were able to engage the primary literature.

Two of our student researchers went on to present their findings at regional and national conferences and won prizes for their posters. One student went on to obtain a Research Experiences for Undergraduates (REU) internship in a prestigious 4-year college. Our preliminary data illustrate the abundant opportunities provided by LAMP technology for students to develop skills in critical thinking and experimental design. Students can optimize multiple parameters, troubleshoot unexpected or discrepant findings, develop comparisons with alternative assays, and test hypotheses with respect to the incidence and distribution of multiple microbial microorganisms. A student response survey data shows that majority of the students had a positive and engaging UR experience (**Figure 7**).

TABLE 3 | Reagent cost comparison between LAMP, PCR, and qPCR.

	Real time	Traditional	LAMP	
	PCR (per reaction)	PCR (per reaction)	(per reaction)	
Enzyme				
(Taq or Bst) SYBR TM Green PCR Master mix 1 ml = \$125 (ThermoFisher) Taq Polymerase	\$2.50	- \$0.47 per Unit	- - \$0.07	
500 Unit (U) = \$236 0.25-2.5 U/50 μl reaction (ThermoFisher) <i>Bst</i> Enzyme 8,000 U/ml = \$70 (New England Biolabs)				
Plates or Strip 300 strips = \$126	\$0.42/strip \$0.0525/well	-	_	
Microfuge tubes (ThermoFisher) 500 = \$22	\$0.044	\$0.044	\$0.044	
PCR tubes 1,000 = \$119 (ThermoFisher)	-	\$0.12	_	
Primers: $(0.1-1 \mu M)$ $100 \mu M-$40$ dNTP's 100 mM $4 \times 250 \mu I = 320 (ThermoFisher)	Forward and Backward \$0.04	Forward and Backward \$0.04 50 µM each 2,000 reactions \$0.16	F3/B3 = \$0.04 LF/FB = \$0.04 FIP/BIP = \$0.08 50 μM each 2,000 reactions \$0.16	
Agarose gel (ThermoFisher) 100 g = \$145	1% gel–\$1.45 \$0.08 per lane	1% gel–\$1.45 \$0.08 per lane	1% gel–\$1.45 \$0.08 per lane	
TOTAL*	\$2.71 [¥]	\$0.91 ⁸	\$0.52 ⁸	

*This is an underestimate as plastic tips/DNA markers/buffers are not calculated. [¥]Most real time PCR reactions are run in triplicate with appropriate internal controls.

⁸Samples can be run in duplicate.

The estimate here is within reported data for the cost per LAMP reaction to be approximately 60–70 cents (1) while most DNA amplification cost between \$1–2 per reaction and \$5–7 per reaction for real time PCR (2).



FIGURE 7 | Survey of student attitudes. An anonymous survey was conducted using Google forms.



Loop-mediated isothermal amplification technology shares with PCR a notable potential to connect URE with important public health issues. Since *E. coli, Salmonella* spp., *Enterococcus* spp., and *S. aureus* spread via contaminated food or water, as well as via infected individuals, environmental detection of these microorganisms can limit the occurrence and magnitude of outbreaks (Curran, 2017). Notably, LAMP has emerged as a relevant technology for the detection of norovirus, *Clostridium perfringens*, and *Campylobacter jejuni*, three of the five most prevalent foodborne pathogens in the United States (Fukuda et al., 2006; Yamazaki et al., 2009; Kaneko et al., 2011), and has recently been used to detect COVID-19 (Kashir and Yaqinuddin, 2020). Students could use LAMP to detect these and other types of microorganisms in the course of developing their own projects. The only additional reagents required would be new primers.

Incorporating authentic research in undergraduate curriculum is challenging due to technical demands and time constraints (Smyth, 2017). Implementing LAMP-based research projects as a CURE is a potential solution. Our data show that both cultures and environmental samples can be effectively processed for LAMP by simple boiling, which should greatly facilitate the design of these experiences in a classroom setting. We intend to implement LAMP-based CUREs in several major and non-major Biology courses using environmental samples. Students will be encouraged to formulate hypotheses as to which environmental sites or site categories are likely to harbor the greatest prevalence of pathogenic microorganisms. Faculty mentors will assist by engaging students in learning about microbes in the context of urban ecology and public health. Pre/post surveys will be used to assess impact on student understanding of scientific method and microbial biology, as well as attitudes toward potential STEM careers.

Finally, a unique feature of LAMP is the potential for its use in distance learning, and more broadly for both studentand community-based research activity, presenting an unusual opportunity for the expansion of public engagement in science as well as interaction with the research community. This potential developmental sequence is summarized in Figure 8. In the context of distance learning, LAMP is uniquely suited to at-home use in a manner that parallels on-site procedures. All components are stable, including the Bst polymerase, which displays remarkable stability (Meridian Bioscience). Any household with an oven, a pot and a thermometer can establish the necessary constant-temperature incubation conditions, and colorimetric readout is readily accomplished. The boiling method for DNA prep can be readily adapted for at-home implementation: inexpensive oven-safe evaporating dishes can be used for sample concentration, substituting for centrifugation. Non-specific products are expected to be rare due to the requirement for hybridization at six independent sequences in the target. Furthermore, it is likely that paper strip-based readout methods will become available that will facilitate semiquantitative analysis (Hongwarittorrn et al., 2017). Quantitation by such methods, as well as by dilution, will allow at-home students to pursue expanded LAMP-based lab activities, including error analysis and protocol variations, which take advantage of the time and flexibility afforded by flipped lab designs. Similarly, at-home CUREs offer expanded possibilities in comparison to on-site research, as students can develop research questions and collect samples in a manner that is tailored to the unique as well as shared aspects of their individual locations. Continual access to the home "laboratory" is likely to encourage students to generate additional research projects in an open-ended manner, providing an unusual opportunity for training in project design at the undergraduate level.

Moreover, the possibilities for at-home LAMP are not limited to the student body, which may serve as a vanguard for the encouragement of similar scientific research activity in communities. Therein potentially lies a unique opportunity for the development of the public culture of science. For example, while it is becoming increasingly common for scientists to crowdsource sample acquisition from the lay public, LAMP is perhaps the unique technology that may permit lay crowdsourcing, not only of samples, but also of methodological data. Lay LAMP-tinkerers may make a contribution to the professional research community, if their numbers outweigh their lack of scientific sophistication. Scientists may contribute to and help guide this process by offering research challenges to such lay communities. Communities of citizen scientists can potentially organize projects of their own, and they may also be a source of sociocultural data for the professional study of the culture of science (Figure 8). By engaging the public this way, we anticipate that this may increase understanding and engagement with science as well as broadening access to and increasing inclusiveness in the process of science.

In summary, incorporation of LAMP in URE has many benefits, including low cost, speed, ease of training, and the ability to engage undergraduates in meaningful UREs. Moreover, beyond these technical advantages, LAMP has unusual promise as a technology for expanding student and community experience in authentic scientific inquiry. UR has become a hallmark national trend as a component of HIP pedagogy. We have been able to observe the value of UR to undergraduate education on a personal level, noting the growth in self-confidence, independence and communication skills in our students as they progress through the program. Further development of LAMPbased approaches to UR will allow students to gain first-hand knowledge of STEM careers, and more broadly to apply newly acquired skills and experience to their further education and professional development.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/**Supplementary Material**, further inquiries can be directed to the corresponding author/s.

AUTHOR CONTRIBUTIONS

All authors contributed to the design of the experiments and agreed to working on different projects. AN planned the project, supervised, and wrote an initial draft. AO revised and rewrote the manuscript, analyzed data, and added a figure. DS and MT contributed to the revision. AN, AO, and MT designed and implemented IRB-approved student survey. CP and MK provided discussion and strategic approach to LAMP research regarding *Enterococcus* and *Salmonella*, respectively. All the students (KP, GL, DM, OB, MF, JS, WK, EC, YK, TN, MF, JE, and MM) performed experiments, data analysis and participated in manuscript preparation. All authors critically discussed and contributed to the final version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fmicb. 2020.603381/full#supplementary-material

Supplementary Figure 1 | Comparison of DNA amplification using PCR (A) or LAMP (B). PCR requires the use of a thermocycler and electrophoresis equipment (A). In LAMP, self-priming obviates the need for a thermocycler, and colorimetric readout provides an alternative to electrophoresis (B).

Supplementary Figure 2 | Schematic diagram of LAMP. *Part 1.* Template DNA with primer target sequences indicated (F3c, F2c, and F1c). *Part 2.* Primer FIP binds to target sequence F2c, with a 5' overhang containing F1c (top panel). Primer F3 is used to unzip and release strand A2 (middle panel), which then forms a loop at its 5' end via F1c-F1 complementarity (bottom panel). *Part 3.* The 3' end of strand A2 is targeted by primer B1P, followed by unzipping with primer B3 to release strand A4, which can forms loops at both 5' and 3' ends. *Part 4.* The resulting dumbbell (strand A4) is comparable to a similar dumbbell (strand B4) formed by a corresponding series of events (not shown) beginning with targeting of B2c in strand B (*Part 1). Part 5.* Loop amplification yields concatemers of target sequences, beginning either with strand A4 (*Part 5a*) or with strand B4 (*Part 5b*).

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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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