

Application of Loop-Mediated Isothermal Amplification combined with colorimetric and lateral flow dipstick visualization as the potential point-of-care testing for diphtheria

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Research article

Keywords: LAMP; POCT; diphtheria; *Corynebacterium diphtheriae*; colorimetric detection; lateral flow dipstick

Posted Date: September 25th, 2019

DOI: <https://doi.org/10.21203/rs.2.14953/v1>

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Version of Record: A version of this preprint was published at BMC Infectious Diseases on April 25th, 2020. See the published version at <https://doi.org/10.1186/s12879-020-05037-z>.

Abstract

Background Diphtheria outbreaks occurred in endemic areas and imported and indigenous cases are reported in UE/EEA. Because of the high infectiveness and severity of the disease, early and accurate diagnosis of each suspected case is essential for the treatment and management of the case and close contacts. The aim of the study was to establish simple and rapid testing methods based on Loop-Mediated Isothermal Amplification (LAMP) assay for the detection of *Corynebacterium diphtheriae* and differentiation between toxigenic and non-toxigenic strains. **Methods** *Corynebacterium diphtheriae* and *Corynebacterium ulcerans* isolates from the National Institute of Public Health-National Institute of Hygiene collection were used for the development of LAMP assay for the diagnosis of diphtheria and nontoxigenic *C. diphtheriae* infections. Various colorimetric methods for visualization of results were investigated. Sensitivity and specificity of the assay were examined using a collection of DNA samples from various gram-positive and gram-negative bacteria. **Results** The LAMP assay for *tox* and *dtxR* genes was developed. The sensitivity and specificity of the assay were calculated as 100%. The detection limit was estimated as 1.42 pg/ μ l concentration of DNA template when the reaction was conducted for 60 min. However, the detection limit was lowered 10 times for every 10 minutes of reduction in the time of incubation during the reaction. Positive results were successfully detected colorimetrically using hydroxynaphthol blue, calcein, QuantiFluor, and lateral flow Milenia HybriDetect dipsticks. **Conclusion** The assay developed in the study might be applied for point-of-care testing of diphtheria and other *C. diphtheriae* infections. It is highly sensitive, specific, inexpensive, easy to use, and suitable for low-resource settings.

Background

Diphtheria is an acute, potentially lethal infectious disease of humans caused by diphtheria toxin-producing strains of *Corynebacterium diphtheriae*, *Corynebacterium ulcerans*, and *Corynebacterium pseudotuberculosis*. It is a highly infectious disease. The infection can be transmitted through contact with infected persons and objects that are touched by them. Depending on the anatomic site that is affected by the disease, it could be respiratory or cutaneous diphtheria. Rarely other sites can also be affected such as eye, ear, and vulva. Diphtheria toxin absorbed from the mucosal or cutaneous lesions causes toxic damage to organs such as the myocardium, kidneys and nervous system. In respiratory diphtheria cases, formed pseudomembranes can cause obstruction in the airways [1].

After the introduction of vaccination against diphtheria in the 1940s, the infections caused by toxigenic corynebacteria seemed to be well controlled in developed countries. However, infections recorded during the last several years point at *C. ulcerans* and *C. diphtheriae* as reemerging human pathogens. According to ECDC data, the number of confirmed diphtheria cases in EU/EEA increased over three times from 2011 to 2015 [2]. Domestic pets and other animals have been described as novel reservoirs and sources of diphtheria infection [3, 4, 5]. Moreover, diphtheria is endemic in many countries in Asia, the South Pacific, the Middle East, and Eastern Europe and in Haiti and the Dominican Republic; outbreaks in Indonesia, Thailand, Laos, South Africa, Sudan, and Pakistan have occurred since 2011 [6]. According to WHO data, 28,358 cases of diphtheria were recorded between 2012 and 2016 worldwide. In the period 2012–2016, India had the largest total number of reported cases each year, with a 5-year total of 17,497 cases, followed by Indonesia and Madagascar with 2739 and 4492 reported cases, respectively [7]. In addition, the diphtheria cases were described in asylum seekers in Europe [8, 9].

Because of the high infectiveness and severity of the disease, early and accurate diagnosis of each suspected case is essential for the treatment and management of the case and close contacts. Rapid microbiological tests are of high value because clinical diagnosis is not easy and might be confused with other causes, such as streptococcal sore throat or tonsillitis [10]. Misdiagnosis is the high risk particularly in countries where the diphtheria is uncommon. Point-of-care diphtheria testing is especially important in refugee camps and developing countries, where access to medical laboratories is extremely limited as well as in the investigation of an infection source.

Methods

Bacterial strains and DNA extraction

Nine toxigenic and 31 nontoxigenic *Corynebacterium* strains were used in the study. The toxigenic strains included 5 *C. diphtheriae* clinical isolates, one *C. ulcerans* clinical isolate, and reference *C. diphtheriae* strains such as PW8, NCTC 10648, and NCTC 3984. The nontoxigenic strains included 30 *C. diphtheriae* clinical isolates and the reference *C. diphtheriae* strain NCTC 10356. Additionally, other bacterial species that might be present in the respiratory tract, such as *Streptococcus salivarius*, *Streptococcus pneumoniae*, *Streptococcus pyogenes*, *Streptococcus epidermidis*, *Haemophilus influenzae*, *Moraxella catarrhalis*, *Staphylococcus aureus*, *Corynebacterium pseudodiphtheriticum*, *Pseudomonas aeruginosa*, *Escherichia coli*, and *Klebsiella pneumoniae*, were included in the study.

DNA extraction was performed for 24 h bacterial cultures on the medium appropriate for the bacterial species by using DNeasy Blood and Tissue Kit (Qiagen, Germany) according to the manufacturer's instructions.

Loop-Mediated Isothermal Amplification (LAMP)

LAMP primer sets for the detection of *tox* gene coding diphtheria toxin and *dtxR* gene coding global regulator were designed by using LAMP designer software PrimerExplorer V4 (<https://primerexplorer.jp/e/>). Each LAMP primer set included two outer (F3, B3), two inner (FIB, BIP), and two loop primers (LF, LB). The sequences of the oligonucleotide primer sets used in the study are presented in Table 1. For the detection of amplified products using the lateral flow dipsticks, the FIB and BIP primers were labeled with biotin and fluorescein isothiocyanate (FITC), respectively, at the 5' end. For the colorimetric detection of amplified products, unmodified primers were used. Modified primers were obtained from Metabion (Germany) and unmodified primers were obtained from Genomed (Poland). LAMP was carried out in a final reaction volume of 25 μ l. The concentration of primers in the reaction mixture was optimized for each target individually. Finally, the reaction mixture for both targets contained 0.8 μ M of FIB and BIP primers each, 0.2 μ M of F3 and B3 primers each, 0.4 μ M of LF and LB primers each, 1 \times reaction buffer containing 20 mM Tris-HCl, 50 mM KCl, 10 mM (NH₄)₂SO₄, 2 mM MgSO₄, 0.1% Tween 20 (New England Biolabs, USA), 0.2 mM dNTP (Sigma-Aldrich, USA), 0.2 M betaine (Sigma-Aldrich, USA), 8 units Bst 2.0 DNA Polymerase (New England Biolabs, USA), and 2 μ l sample DNA. The reaction was optimized at the temperature ranging from 62°C to 70°C and conducted for 60 min. During the optimization step, the results of the reaction were analyzed using agarose gel electrophoresis.

Detection of product amplification with the lateral flow dipsticks

Milenia HybriDetect dipsticks (Milenia Biotec, Germany) were used for the detection of the amplified products labeled with biotin and FITC. Ten microliters of the reaction mixture were pipetted directly on the sample application area on the dipstick. Then, the dipstick was placed with the same application area into 100 μ l of HybriDetect assay buffer and incubated for 5–15 min in an upright position. The results were regarded as positive when two bands were visible (a control band and a test band) or as negative when only a control band was visible.

Colorimetric detection of amplified products

For the colorimetric detection of amplified products, 5 indicators were used comparatively: neutral red, phenol red, hydroxynaphthol blue (HNB), calcein and QuantiFluor. Neutral red and phenol red are pH indicators. They are added to the pre-reaction solution. The progress of LAMP reaction is related to lowering of the solution pH, which can be observed directly as color change of faint orange to pink (neutral red) or red to yellow (phenol red) [11]. Hydroxynaphthol blue and calcein are metal ion indicators. They are also added to the pre-reaction solution. When Mg²⁺ ion concentration decreases in the progress of LAMP reaction, the color change of the indicators can be observed directly [12]. The color shift is violet to blue for HNB and orange to fluorescent green for calcein. QuantiFluor is a DNA intercalating dye. It is added to the solution after the reaction is completed. When the LAMP reaction is positive, a color change of orange to fluorescent yellow is observed under ambient light condition.

Neutral red (Sigma-Aldrich, USA) and phenol red (Sigma-Aldrich, USA) were dissolved in deionized water or 1 M NaOH, respectively, at 50 mM to prepare a stock solution and diluted to 2.5 mM. For the optimization of the concentration of the indicators in the reaction solution, the following final concentrations were tested: 0.2, 0.15, 0.1, 0.05, and 0.01 mM. HNB was dissolved in deionized water at 50mM to prepare a stock solution. Then, the solution was diluted and tested in the LAMP reaction at the following final concentrations: 1, 0.5, 0.32, 0.25, 0.16, 0.125, 0.08, and 0.04 mM. The calcein stock solution consisted of 0.5 mM calcein (Novazym, Poland) and 10 mM MnCl₂ (Sigma-Aldrich, USA) in deionized water. To select an optimal concentration, the following volumes of the stock solution were added to the reaction solution: 0.25, 0.5, 1, 1.5, and 2 μ l. The amount of QuantiFluor (Promega, Germany) in the post-reaction solution was optimized by the addition of the following volumes of the dye: 2, 1, and 0.5 μ l.

Determination of specificity, sensitivity, detection limit, and minimal reaction time

Specificity and sensitivity of the LAMP were investigated using abovementioned bacterial species that can be present in respiratory tracts. The sensitivity was calculated as follows: $A/(A + C) \times 100\%$, and the specificity was calculated as follows: $D/(B + D) \times 100\%$, where A is the number of true positive results, B is the number of false-positive results, C is the number of false-negative results, and D is the number of true negative results.

The limit of detection was investigated using 10-fold serial dilutions of the total genomic DNA.

To determine required minimal LAMP reaction time, we examined the results of the reactions for *tox* and *dtxR* markers after 10, 20, 30, 40, 50, and 60 min of incubation using 10-fold serial dilutions of the total genomic DNA as a reaction template.

Results

The species-specific *dtxR* gene present in all *C. diphtheriae* strains and the *tox* gene present only in potentially toxigenic *C. diphtheriae*, *C. ulcerans*, and *C. pseudotuberculosis* strains were selected as target genes for designing the LAMP primers. Initially, three sets of primers for each of the genetic markers investigated were designed but only the sets presented in Table 1 did not yield false-positive results and therefore were selected for the study. The concentration of each of the primer as well as other reagents in the reaction mixture was optimized. Labeling of the primers FIB and BIP with biotin and FITC did not influence the amplification reaction, as it was assessed based on agarose gel electrophoresis results. The efficiency of the LAMP reaction was comparable in the temperature ranging from 62°C to 70°C (Figure 1). For the study, we selected 65°C as recommended by the manufacturer of the Bst 2.0 DNA polymerase.

We could detect positive LAMP reactions with the naked eye using HNB, calcein, QuantiFluor, and Milenia HybriDetect dipsticks. The positive reaction was clearly visible when the used HNB concentration was 0.125, 0.16, and 0.25 mM. For further studies, we selected the concentration 0.16 mM of HNB. The optimal amounts of calcein and QuantiFluor per reaction were 0.5 and 2 µl, respectively. By using Milenia HybriDetect dipsticks, we observed atypical results for samples with a high concentration of DNA. According to manufacturer's instructions, two color bands should be visible on the dipstick for positive results: test band and control band, whereas only control band should be visible for negative results. However, we observed that when the concentration of amplicons was high, the control band was not visible (Figure 2). This issue was overcome by the dilution of the amplified product. We could not detect positive LAMP reaction when the neutral red and phenol red were used. It was probably because the pH changes during the reaction were very subtle.

The sensitivity and specificity of the LAMP reaction for *tox* and *dtxR* markers were comparable using HNB, calcein, and QuantiFluor, as well as Milenia HybriDetect dipsticks, and both were calculated as 100% (Table 2). The detection limit was also comparable for both genetic markers and all product detection methods and estimated as 1.42 pg/µl concentration of DNA template, which means 2.84 pg of DNA in 25 µl of the reaction mixture, when the reaction was conducted for 60 min. However, the detection limit lowered 10 times for every 10 minutes of reduction in the time of incubation during the reaction (Table 3, Figure 3).

Discussion

Diphtheria is a vaccine-preventable disease. Currently, the diphtheria vaccination coverage varied from 42% in some developing countries to 99% in some developed countries [13]. Diphtheria outbreaks occur in endemic countries, and diphtheria cases are reported every year in UE/EEA. According to ECDC data, 216 diphtheria cases were reported in UE/EEA in the period 2012–2016. Most of them were imported from endemic geographical areas and some were indigenous cases [14]. Epidemiological data on diphtheria and a growing problem of nontoxigenic *C. diphtheriae* invasive infections [15, 16, 17] have revealed the need for point-of-care testing (POCT) technology for the detection of *C. diphtheriae* in carriers, suspected cases, and contacted persons. Such POCT technology would be of great value especially in endemic regions of the disease, where access to health care is limited, and in refugee camps, to timely start appropriate treatment and further prevent the spread of the *C. diphtheriae* and the outbreak. Our study aimed to establish simple and rapid testing methods based on LAMP assay for the detection of *C. diphtheriae* and differentiation between toxigenic and non-toxigenic strains. Additionally, we compared various methods for visualization of amplified products. The developed method was highly sensitive and specific and showed a very low detection limit. It was reported by other researchers that LAMP detection limit is lower than polymerase chain reaction (PCR) [18]. However, we found that the incubation time necessary to obtain positive results depends on the amount of the target DNA in the sample. The methods of visualization the results, including the use of HNB, QuantiFluor, calcein, and Milenia HybriDetect dipsticks, did not influence the detection limit, but colorimetric detection using HNB and calcein, which were added to the reaction mixture before incubation, are superior to QuantiFluor and Milenia HybriDetect dipsticks because they enable faster detection of positive reaction in the real-time mode, and no additional handling after reaction is needed. The opening tubes after the reaction is associated with an increased risk of contamination of other subsequent LAMP reaction solutions.

The LAMP reaction temperature ranging from 62°C to 70°C, as revealed in our study, shows that the heating device used does not have to be very precise. Hatano et al. [19] proposed the conduction of LAMP using a disposable pocket warmer placed in a Styrofoam box. It makes the LAMP assay independent from any electric power and therefore applicable in sites where electricity infrastructures are inadequate, such as undeveloped areas. LaBarre et al. [20] developed the non-instrumented nucleic acid amplification heater suitable for isothermal amplification methods, where heat is generated by an exothermic reaction of CaO with water.

One of the disadvantages of most molecular methods is the requirement for storage conditions of reagents, such as polymerases, which usually have to be kept in freezing. However, a lyophilized mastermix containing all reagents required for LAMP assay was developed for the detection of some other pathogens [20, 21]. Furthermore, LAMP reagents are commercially available in a dry format currently, which can be stored at room temperature.

At the stage of development of the potential point-of-care test for diphtheria, we used DNA samples. However, according to Yan et al. [22] and Dugan et al. [23], the LAMP assay can be conducted directly from the bacterial colony, as high temperature causes the leak of bacterial cells, after which a high amount of DNA is released. Therefore, our test might be validated to be used for clinical samples directly. It was confirmed that LAMP assay was less affected by various components of clinical samples compared to other molecular methods, such as PCR [22, 24], and therefore, the DNA samples do not have to be purified perfectly.

Conclusions

The developed LAMP for diphtheria diagnostics might be a valuable tool for outbreak investigations, especially in endemic areas, as well as for rapid screening of travelers coming from diphtheria-endemic regions. It can also be used for the examination of carriers and diphtheria contact persons. The selection of *dtxR* gene, apart from *tox* gene, as a diagnostic marker enables also the detection of non-toxicogenic *C. diphtheriae* strains in tested samples, which is important in countries where nontoxicogenic *C. diphtheriae* invasive infections are reported [15, 16].

The assay developed in the study has the potential to be integrated into a diagnostic mobile device for POCT of diphtheria and other *C. diphtheriae* infections, which will be highly sensitive, specific, inexpensive, easy to use, and applicable in low-resource settings. Although our study revealed high specificity and sensitivity and low limit detection of the assay, it should be tested using real clinical samples directly.

Abbreviations

LAMP - Loop-Mediated Isothermal Amplification

HNB - hydroxynaphthol blue

POCT - point-of-care testing

UE/EEA - European Union/European Economic Area

Declarations

Ethics approval and consent to participate

Ethical approval was not required for the use of clinical samples because all the investigated isolates were taken as a part of standard care (diagnostic purposes). The samples were not collected for research purposes.

Consent for publication

Not applicable.

Availability of data and material

All data generated or analyzed during this study are included in this published article.

Competing interests

The authors declare that they have no competing interests.

Acknowledgments

Not Applicable

Funding

This study was supported by the National Institute of Public Health – National Institute of Hygiene internal-grant (no. 1/EM/2018). The funder had no role in the study design, data collection, analysis and interpretation and preparation of the manuscript.

Authors' contributions

AAZ designed the study, optimized the reaction conditions and prepared the manuscript. AW and UC conducted sensitivity and specificity tests using HNB and determined limit of detection, KB conducted sensitivity and specificity tests using calcein and participated in the determination of detection limit; KF conducted sensitivity and specificity tests using Milenia HybriDetect; EM, MP, and KW conducted sensitivity and specificity tests using QuantiFluor; KKG designed primers. All authors participated in the analysis of results. All authors read and approved the final manuscript. All authors gave final approval of the version to be published and are publicly responsible for its contents.

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Tables

Table 1. LAMP primers used in the study.

Target gene	Name	Sequence (5' → 3')
<i>tox</i>	LF-toxIII	GCATAGTTAGCCCCAGCGAAT
	LB-toxIII	ACTTCCTGGTATCGGTAGCGT
	F3-toxIII	CGGCATTAGAGCATCCTG
	B3-toxIII	CTAGCTCTCCTACCAATGGA
	FIP-toxIII	CGCAACGTTTACTGCCCATTTTCTTACTGGGACCAATCCTGT
	BIP-toxIII	AAGACAACCTGCTGCTCTTTTTTTCGATATTGTGGTGAACGGCAC
<i>dtxR</i>	LF-dtxRIII	TCGTCACCTCATAACGTGTTCC
	LB-dtxRIII	CGGCGTAGGCAATTCTGA
	F3-dtxRIII	AACATCGCTTAGCTGAGC
	B3-dtxRIII	CGTTAATCTGAACAATGCGTAC
	FIP-dtxRIII	TTCACGAGCCTGCGTTCTTTTAAAAGTTTACGATGAAGCCTG
	BIP-dtxRIII	CAATTCCAGGTCTCGACTTTTTGAACTTCAATAACGCGAGTTCCG

Table 2. Results of the LAMP for *tox* gene and *dtxR* gene for various DNA samples.

Bacterial species (number of strains tested)	LAMP results	
	<i>tox</i>	<i>dtxR</i>
Toxigenic <i>Corynebacterium diphtheriae</i> (31)	+	+
Non-toxigenic <i>Corynebacterium diphtheriae</i> (8)	-	+
Toxigenic <i>Corynebacterium ulcerans</i> (1)	+	-
<i>Streptococcus salivarius</i> (1)	-	-
<i>Streptococcus pneumoniae</i> (1)	-	-
<i>Streptococcus pyogenes</i> (1)	-	-
<i>Streptococcus epidermidis</i> (1)	-	-
<i>Haemophilus influenzae</i> (1)	-	-
<i>Moraxella catharalis</i> (1)	-	-
<i>Staphylococcus aureus</i> (1)	-	-
<i>Corynebacterium pseudodiphtheriticum</i> (1)	-	-
<i>Pseudomonas aeruginosa</i> (1)	-	-
<i>Escherichia coli</i> (1)	-	-
<i>Klebsiella pneumoniae</i> (1)	-	-

Table 3. Results of examination the minimal incubation time required for the LAMP assay.

Incubation time (min)	DNA dilution (used as a template)					Negative control
	1.42 ng/μl	142 pg/μl	14.2 pg/μl	1.42 pg/μl	142 fg/μl	
10	-	-	-	-	-	-
20	-	-	-	-	-	-
30	+	-	-	-	-	-
40	+	+	-	-	-	-
50	+	+	+	-	-	-
60	+	+	+	+	-	-

Figures

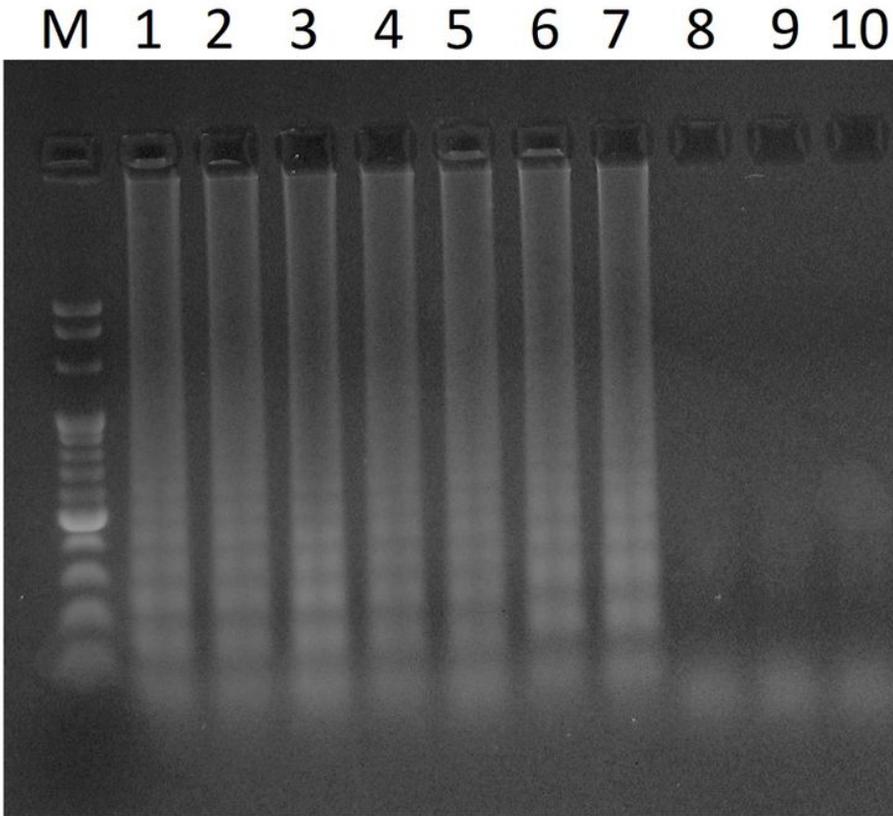


Figure 1

Agarose gel electrophoresis of products of the LAMP reaction for tox gene conducted for 60 min at various temperatures. M—Molecular Ladder, lane 1—incubation at 62°C, lane 2—incubation at 64°C, lane 3—incubation at 65°C, lane 4—incubation at 66°C, lane 5—incubation at 67°C, lane 6—incubation at 68°C, lane 7—incubation at 70°C, and lanes 8–10—negative controls.



Figure 2

Visualization of LAMP for dtxR gene using lateral flow dipsticks. A—serial dilutions of the amplified product; B—serial dilutions of the DNA sample.

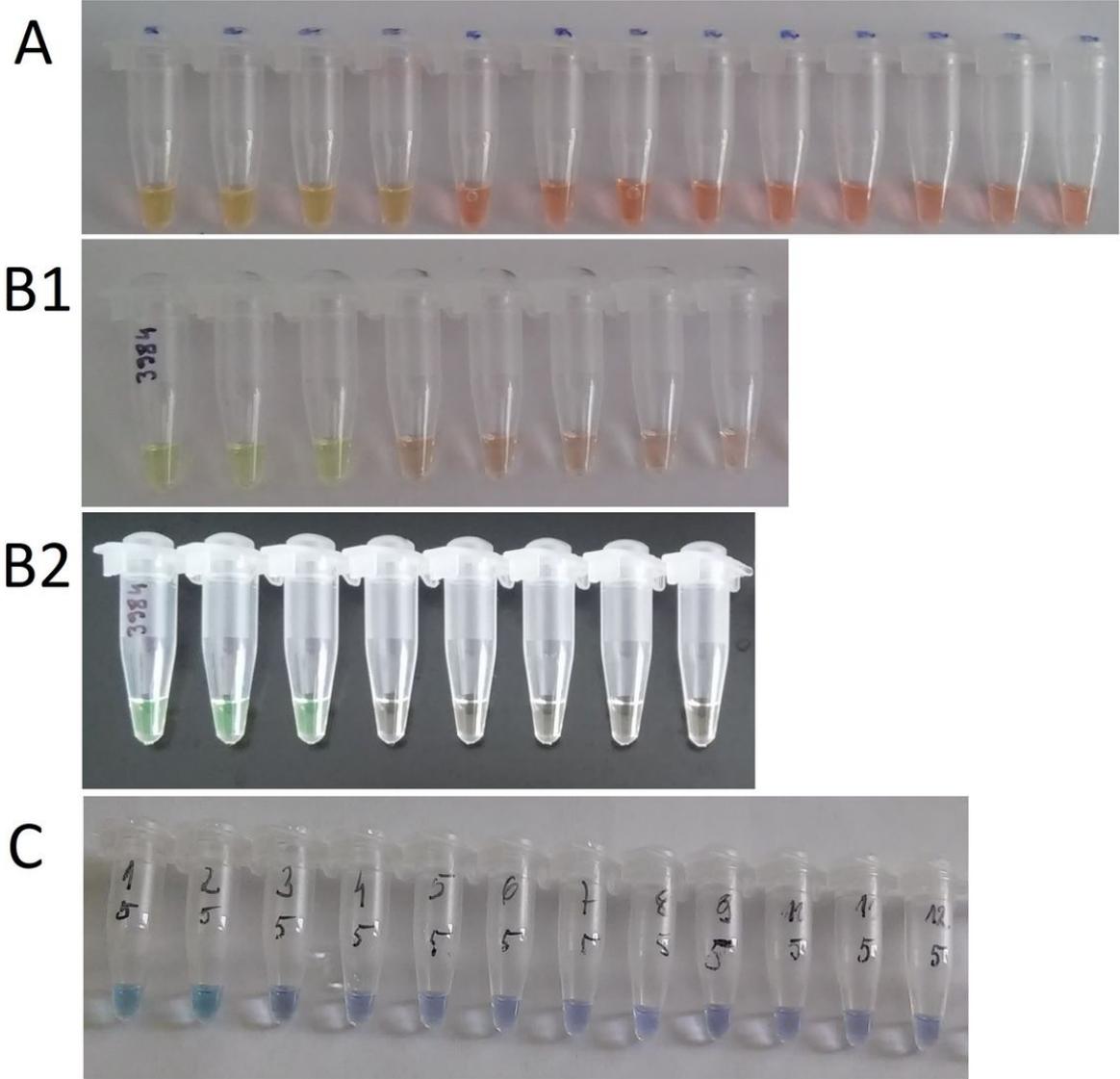


Figure 3

Results of the LAMP detection limit for tox gene. From the left to the right, 10-fold serial dilutions of the DNA samples. A—60 min of incubation, detection using QuantiFluor; B—50 min of incubation, detection using calcein (B1—white background, B2—black background); C—40 min of incubation, detection using HNB.