# Development and evaluation of a rapid visual loop-mediated isothermal amplification assay for the *tcdA* gene in

## Clostridioides difficile detection

4 5

1

2

3

- $6 \qquad \text{Minyi Lin}^{1,\#}, \text{Pu Wang}^{3,\#}, \text{Bingyun Lu}^2, \text{Ming Jin}^2 \text{ , Jiasheng Tan}^4, \text{Wei Liu}^5 \text{ , Jing Yuan}^5, \text{Xiaomou}$
- 7 Peng<sup>1,\*</sup> and Ye Chen<sup>2,\*</sup>

8

- 9 Department of Infectious Diseases, The Fifth Affiliated Hospital, Sun Yat-Sen University, Zhuhai,
- 10 Guangdong, China
- 11 <sup>2</sup> Integrative Microecology Center, Shenzhen Key Laboratory of Gastrointestinal Microbiota and Disease,
- 12 Shenzhen Clinical Research Center for Digestive Disease, Shenzhen Technology Research Center of Gut
- 13 Microbiota Transplantation, Shenzhen Hospital, Southern Medical University, Shenzhen, Guangdong,
- 14 China
- 15 <sup>3</sup> Department of Gastroenterology, Guangdong Provincial Key Laboratory of Gastroenterology, State Key
- 16 Laboratory of Organ Failure Research, Nanfang Hospital, Southern Medical University, Guangzhou,
- 17 Guangdong, China
- 18 <sup>4</sup>Department of Gastroenterology, SongShan Lake Central Hospital of Dongguan City, Dongguan,
- 19 Guangdong, China
- <sup>5</sup> Institute of Disease Control and Prevention, Academy of Military Medical Sciences, Beijing, China
  - <sup>#</sup> These authors contributed equally to this work.

21 22

- 23 Corresponding Author:
- 24 XiaoMou Peng
- 25 52 East Meihua Road, Zhuhai, Guangdong, 519000, China
- 26 Email address: pengxmou@mail.sysu.edu.cn
- 27 Ye Chen,
- 28 1333 New Lake Road, Shenzhen, Guangdong, 518100, China
- 29 Email address: yechen@smu.edu.cn

30 31

32 33 34

35 36 37

38 39 40

41 42

Abstract

**Background.** The *tcdA* gene is of great significance to the codes for an important toxin earrying status of produced by *Clostridioides difficile* (*C. difficile*), but there is currently no simple and cost-effective method of detecting it. This paper establishes and validates a rapid and visual loop-mediated isothermal amplification (LAMP) assay for the detection of the *tcdA* gene of *C. difficile*.

**Methods**. Three sets of primers were designed and optimized to <u>target amplify</u> the *tcdA* gene in *C*. *difficile* using a LAMP assay. The sensitivity and specificity of LAMP primers were detected by The *tcdA* gene of *C*. *difficile* was detected in 164 stool specimens using both LAMP and polymerase chain reaction (PCR). Positive and negative results were distinguished using real-time monitoring of turbidity and chromogenic reaction.

**Results**. At a temperature of 66 °C, the target DNA was successfully amplified with all three sets of LAMP primers, and visualized within 60 minutes, for a *C. difficile strain that carries the tcdA gene*. Under the same conditions, The results of the target DNA was not amplified with the all/two/one set(s) of LAMP primers detection of for 26 pathogenic bacterial strains without the that do not carry the tcdA gene were negative, indicating that the primers were extremely specific. Moreover, the detection limit of LAMP was  $\frac{20.721}{20.721}$  pg/µl, which was 10 times more sensitive than that of conventional PCR. Finally, the The detection rate of tcdA in 164 stool specimens, using the LAMP method, was 17% (28/164), significantly higher than the 10% (16/164) detection rate of the PCR method ( $X^2=47$ , p < 0.01). Conclusion. In this study, the LAMP method proved to be is an effective technique for the rapid and visual detection of the tcdA gene of *C. difficile*, demonstrating and shows potential advantages over PCR

visual detection of the *tcdA* gene of *C. difficile*, demonstrating and shows potential advantages over PCR in terms of speed, simplicity, and sensitivity. The *tcdA*-LAMP assay is particularly suitable for medical diagnostic environments with limited resources and is a promising diagnostic strategy for the screening and detection of *C. difficile* infection in populations at high risk. Future research will need to further validate the applicability of the LAMP technique across various experimental and clinical settings and address any technical challenges that may arise during its broader implementation.

Introduction

Clostridioides difficile (C. difficile) is a serious problem in health care institutions the major cause of antibiotic-associated diarrhea worldwide (Rodríguez et al., 2020). Although the estimated national burden of C. difficile infection (CDI)<sub>2</sub> and associated hospitalizations<sub>2</sub> decreased from 2011 through 2017 in 10 USA states (California, Colorado, Connecticut, Georgia, Maryland, Minnesota, New Mexico, New York, Oregon, and Tennessee), the number of C. difficile cases in the USA remained as high as 15,512 in 2017, with an estimated total national burden of 462,100 cases and an estimated incidence of 144 cases per 100,000 population (Guh et al., 2020). In Shandong and Zhejiang provinces, China, tertiary hospitals reported a consistent 14% incidence of CDI (72/504) among hospitalized patients with suspected CDI in Shandong and 14% (115/804) among acute gastroenteritis outpatients in a Zhejiang pediatric hospital (Luo et al., 2018; Shuai et al., 2020). These findings are consistent with other studies from mainland China, which report a 14% crude incidence of toxigenic C. difficile in diarrheal patients (Tang et al., 2016).

With the global prevalence of CDI, specific and sensitive methods for identifying *C. difficile* are needed. The primary virulence factors of *C. difficile* are two structurally similar toxins, denoted as toxin A and toxin B (tedA and tedB, which are encoded by the tedA and tedB genes respectively; (Kuehne et al.,

**Commented [LMG1]:** Need to describe methods (in silico analysis, number of positive and negative control *strains...*)

Commented [LMG2R1]: It sounds like by testing for amplification with non-C. diff bacteria and then comparing to PCR for stool samples of patients that showed symptoms of CDI

Commented [LMG3]: It is not at all clear in if the results presented in Abstract apply to 1, 2 or 3 sets of the primers mentioned in Methods

**Commented [LMG4]:** Of the three primer sets, how many were positive?

**Commented [LMG5]:** Use reasonable significant figures

Commented [LMG6]: Focus on your work, not the future

**Commented [LMG7]:** New paragraph needed to state the unmet need

2011). Most cases of CDI are attributed to strains expressing both toxins A and B (A+B+) (ref). Although the toxin A-B+ strain is uncommon, it can also cause disease and has been relevant to previous outbreaks of CDI (Alfa et al., 2000; Kuijper et al., 2001; Drudy et al., 2007). It was previously believed that the Earlier reports indicated that toxin A+B- strain only rarely caused human diseases (Rupnik, 2008). As a result, numerous studies have emphasized the importance of toxin B in the pathogenesis of CDI, while downplaying the importance of toxin A. However, a study recently reported the discovery of clinical pathogenic C. difficile strains that produce high levels of toxin A but minimal or no toxin B, indicating that toxin A alone can cause CDI (Lin et al., 2020). This pattern of toxin production, observed in more than 5% of isolates, is consistently found both *in vitro* and *in vivo* in humans and mice (Lin et al., 2020). Furthermore, the production of either toxin A or toxin B by these isolates is sufficient to induce the full spectrum of CDI symptoms (Drudy et al., 2007; Freeman et al., 2010). Additionally, it has been demonstrated in studies that both toxins A and B can independently cause disease in animal models (Kuehne et al., 2011; Kuehne et al., 2010). These studies indicate that greater attention should be paid to toxin A because of its pathogenicity. Current laboratory tests for the diagnosis of toxin A in C. difficile strains include the C. difficile cytotoxin neutralization assay (CCNA), toxigenic culture (TC), toxin A enzyme immunoassay (EIA), glutamate dehydrogenase (GDH) assay, and nucleic acid amplification test (NAAT). Although CCNA and TC remain the current gold standards, their use for routine clinical detection is challenging due to their requirements for harsh culture conditions, involvement of highly technical and complex operations, and time-consuming nature (Shah et al., 2020; Liu et al., 2021). Enzyme immunoassays are specific and rapid, but not sensitive (Nicholson & Donskey, 2023). The glutamate dehydrogenase assay is sensitive and rapid, but it has some disadvantages such as eross reactivitycrossreactivity, poor specificity, and a high false positive rate (Bartlett, 2010; Crobach et al., 2016). Previous studies have developed and evaluated ways of Assays for detecting the tcdA gene encoding toxin A by NAAT including include Polymerase Chain Reaction (PCR; Kim et al., 2022), multiplex-PCR (Moosavian et al., 2022), quantitative real-time PCR (Brennhofer et al., 2022), and multiplex real-time PCR (Novakova et al., 2021). Despite the specificity and sensitivity of these diagnostic methods, their suitability for rapid detection in primary hospitals and on-site detection is limited due to their timeconsuming and complex nature, as well as the requirement for expensive equipment. Thus, a rapid, simple, and cost-effective assay is needed to complement current PCR methods for detecting the tcdA gene. Loop-mediated isothermal amplification (LAMP) is a powerful molecular technique for nucleic acid amplification. LAMP, leveragesing the strand displacement activity of Bst DNA polymerase, which facilitates DNA amplification under isothermal conditions (Notomi et al., 2000; Ushikubo, 2004). The high amplification efficiency of this technique, - capable of generating up to 109 copies of target DNA within an hour - underscores its potential in rapid diagnostic applications. The excellent specificity of LAMP is attributed to its use of four (or six) primers, which can identify six (or eight) distinct regions on the target DNA or RNA (Notomi et al., 2000; Parida et al., 2008). Additionally, the detection limit of LAMP surpasses that of PCR, and the results can be visually interpreted without the need for sophisticated equipment. With its high sensitivity and specificity, LAMP has been effectively used to detect various pathogens, including bacteria (Hong-Min et al., 2023), viruses (Nawab et al., 2024), parasites (Chen et al., 2023), and fungi (Badparva et al., 2022), as well as different toxin types (Norén et

87

88

89

90

91

92

93

94

95

96

97

98

99

100

101

102

103

104

105

106

107

108

109

110

111

112

113

114

115

116

117

118

119

120

121

122

123

124

125

126

127

128

129

130

**Commented [LMG8]:** This is an important point, since the assay isjust for tcdA. Do you have a reference?

**Commented [LMG9]:** Do not say "believed," unless earlier reports were not based on the best available data at the time.

Formatted: Font: Italic

Formatted: Font: Italic

**Commented [LMG10]:** Avoid phrases like this. They have no content.

al., 2011; Pancholi et al., 2012), binary toxin genes (Yu et al., 2017), and resistant genes (Lin et al., 2015;

Lin et al., 2022) of *C.difficile*. This study designed three novel sets of LAMP primers and optimized

LAMP for *tcdA* detection. To ascertain the specificity of the *tcdA* primer within the LAMP assay, 26 distinct pathogenic bacterial strains devoid of the *tcdA* gene were analyzed as negative controls. Primer sensitivity was assessed by conducting serial dilutions of *C. difficile* VPI10463 DNA. Finally, the study compared the consistency of LAMP and PCR methods in detecting the *tcdA* gene of *C. difficile* in 164 stool specimens.

137 Materials & Methods

138 Bacterial strains

131

132

133

134

135

136

139

140

141

142

143

144

146

147

148

149

150

151

152

153

154

155

156

157

158

159

160

161

162

A total of 26 pathogenic bacterial strains were selected for to evaluate the specificity of the LAMP detection assay (Table 1). C. difficile VPI10463, which earrying carries the tcdA gene, was used as the positive control. The tcdA gene of VPI10463 the strain showed 100% identity with those of the tcdA gene in previously reported genes ????? which was confirmed by PCR-based sequencing (Fig. S1). To evaluate the specificity of the LAMP assay, 26 pathogenic bacterial strains lacking the tcdA gene were tested.

145 Clinical stool specimens

Fresh stool specimens of suspected CDI inpatients with diarrhea were continuously collected from August 1, 2013 to February 28, 2014 in Nanfang Hospital of Southern Medical University, Guangzhou, China. Inpatients over 18 years old who had received antibiotic or chemotherapy treatments within the past 60 days were included. Stool samples were included of patients who experienced diarrhea within 48 hours of hospitalization, with no less than three episodes of diarrhea within a 24-hour period, and with shapeless stool classified as Bristol types 5-7. The exclusion criteria for this study were as follows: patients who were under the age of 18; patients who had previously been sampled; patients with chronic diarrhea; patients who had used laxatives; patients with various types of infectious diarrhea, such as bacillary dysentery, typhoid fever, food poisoning, and amebic dysentery; patients with intestinal functional diseases, such as irritable bowel syndrome; patients with other types of diarrhea with clear causes unrelated to antibiotics, such as lactose intolerance; and patient samples that did not complete the entire testing process due to instrument or human errors. Of the 197 fresh stool specimens collected from inpatients with suspected CDI presenting diarrhea, 33 were excluded due to duplication (n=23) or inadequate volume and freshness (n=10). All stool specimens were frozen at 80° C until detection. This study was performed in line with the principles of the Declaration of Helsinki. Approval was granted by the Ethics Committee of Fudan University Affiliated Huashan Hospital (committee ethic number: FDEC-2012-014). As a research member unit, Nanfang Hospital was successfully granted an ethics exemption by the Ethics Committee of Southern Medical University.

163164165

166

167

168

169

170 171

172

DNA extraction

To determine the specificity and sensitivity of the LAMP reactions under real conditions, genomic DNA was extracted from *C. difficile* VPI10463 and purified by the Wizard Genomic DNA Purification Kit (Promega, Madison, Wisconsin, USA). The purified genomic DNA was serially diluted in distilled water by a factor of 10, from 207 ng/ul to 0.000207 pg/ul. The concentration of pure genomic DNA before and after dilution was measured using the ND-1000 spectrophotometer (Thermo Fisher Scientific, Inc., Waltham, Massachusetts, USA). Genomic DNA of 26 pathogenic bacterial strains was extracted using a bacterial genomic DNA extraction kit (Tiangeng, Ningbo, China) according to the manufacturer's

Commented [LMG11]: In what? Provide the strain it matched. Also, I suggest you do a microbial genome blast with the sequence KC292122. I found the top 100 hits were all C. diff with > 98% identity.

**Commented [LMG12R11]:** Excluding C. diff genomes, KC292122 did not show significant similarity to genomes of other bacteria

- instructions. Additionally, a stool genome extraction kit (Tiangeng, Ningbo, China) was employed to extract genomic DNA from stool samples. Genomic DNA was stored immediately at -20 °C until use.
- 176 Primer design

175

186

197

209

- 177 Based on the tcdA gene sequence of C. difficile obtained from NCBI GenBank database (GenBank
- 178 accession number: X92982.1), three sets of LAMP primers were designed (Table 2). Further analysis of
- the sequences with the Primer Explorer V4 software [http:primerexplorer.jp/lamp] yielded the outer
- 180 forward primer (F3), outer backward primer (B3), forward inner primer (FIP), and backward inner primer
- 181 (BIP). The FIP and BIP primers recognized both sense and antisense strands and were linked by a four-
- thymidine spacer (TTTT). The two loop primers (LF and LB) were designed to accelerate the
- 183 amplification reaction. To compare the sensitivity and specificity of LAMP and PCR, conventional PCR
- was performed with the NK1 and NK2 primers (Table 2; Kato et al., 1991). All the primers were
- synthesized commercially (Sangon Biotech Co., Ltd., Shanghai, China).
- 187 LAMP reaction
- 188 The LAMP reactions were performed in a 25 µl reaction mixture (DNA amplification kit; Eiken Chemical
- 189 Co., Ltd., Tochigi, Japan) containing the following reagents in the final concentration: 20 mM Tris-HCl
- 190 (pH 8.8), 10 mM (NH4)2SO4, 10 mM KCl, 0.1% Tween-20, 0.8 M betaine, 8 mM MgSO4, 1.4 mM
- 191 deoxynucleoside triphosphate and 8 U Bst DNA polymerase. Each LAMP reaction, using a real-time
- 192 turbidimeter, was composed of 40 pmol FIP and BIP, 20 pmol LB and LF, 5 pmol F3 and B3 primers,
- 193 and 2 µl DNA template. An additional 1 µl of calcein/Mn<sup>2+</sup> complex (Fluorescent Detection Reagent;
- 194 Eiken Chemical Co., Ltd., Tochigi, Japan) was added if direct visual inspection was required. The
- 195 reaction was conducted in reaction tubes (Eiken Chemical Co., Ltd., Tochigi, Japan) within 60 minutes at
- 196 an isothermal temperature of 66  $^{\circ}$ C.
- 198 Detection of LAMP products
- 199 Data were collected as previously described in Liu et al. (2022). Specifically, two different methods,
- 200 chromogenic reaction with calcein/Mn<sup>2+</sup> complex and real-time monitoring of turbidity, were applied to
- 201 detect LAMP products. For direct visual inspection, 1 µl of calcein (fluorescent detection reagent; Eiken
- 202 Chemical Co., Ltd., Tochigi, Japan) was added to 25 µl of reaction mixture in a LAMP tube before the
- 203 LAMP reaction. For a positive reaction, the color changed from orange to green, while a negative reaction
- failed to turn green and remained orange. The color change could be observed by naked eye observation
- 205 under natural light or 365 nm ultraviolet light. For assessing turbidity (Mori et al., 2001), real-time
- 206 amplification was monitored through spectrophotometric analysis by measuring the optical density (λ650
- 207 nm) at 400 nm every 6 s with the aid of a Loopamp real-time turbidimeter (LA-230; Eiken Chemical Co.,
- 208 Ltd., Tochigi, Japan).
- 210 PCR detection
- The PCR conditions used for amplification were described previously (Kato et al., 1991).
- 212 Electrophoresis using a 2% agarose gel (Amresco, Solon, Ohio, USA) containing ethidium bromide was
- 213 applied to analyze the PCR-amplified products. Images were captured using a Bio-Rad Gel Doc EQ
- 214 Imaging System (Bio-Rad, Hercules, CA, USA).

216 Statistical analysis

The required sample size was estimated using Buderer's method (Buderer, 1996), setting the Z-value at 1.96 for the normal distribution and constraining the width of the 95% confidence interval to a maximum of 10%. Previous similar research reported a specificity and sensitivity of 95% for the LAMP assay (Soroka et al., 2021). Given the prevalence of CDI in China is 11% (Wen et al., 2023), the sample size of the study had to be at least 160 to ensure statistical validity. The McNemar test was used to analyze count data, and the Cohen's kappa ( $\kappa$ ) statistic was employed to evaluate the agreement between the LAMP and PCR methodologies. A  $\kappa$  correlation value of 0.40 or below signifies a weak level of agreement, a value ranging from 0.41 to 0.60 reflects moderate agreement, and a value exceeding 0.60 denotes a strong agreement between observations. The specificity, sensitivity, positive predictive value (PPV), and negative predictive value (NPV) of the LAMP and PCR methods were calculated and compared to gene sequencing, respectively. All statistical analyses were performed using SPSS software version 26.0 (IBM

Corp., Armonk, NY, USA). A p-value of <0.05 was considered statistically significant.

Results

Optimal primers for rapid detection of tcdA

All three sets of primers designed herein (Table 2) produced turbidity after 26 minutes (Fig. 1). Primer set tcdA12 showed the fastest amplification (Fig. 1), so tcdA12 primers were selected for optimization. The reaction time of tcdA12 primers with additional loop primers (LB and LF) was less than one-half that of the tcdA8 primers without loop primers.

Appropriate temperature for tcdA LAMP reaction

The *tcdA12* primer set was evaluated across a temperature range of 58 °C to 69 °C at intervals of 1 °C. The ideal temperature range for the *tcdA12* primer set was determined to be 60 °C to 67 °C, with peak amplification efficiency identified at 66 °C (Fig. 2).

Specificity of tcdA LAMP reaction

To estimate the specificity of the LAMP reaction for the tcdA gene, the C. difficile VPI10463 strain with the tcdA gene was used as the positive control. All 26 pathogenic bacterial strains of different species without the tcdA gene and distilled water were designed as the negative controls. As exhibited in Fig. 3A, the increased turbidity curve appeared only when the C. difficile VPI10463 strain with the tcdA gene was used as the template instead of distilled water or other bacterial species. The results were also monitored using a direct visual method (Fig. 3B). Prior to the LAMP reaction, 1  $\mu$ l of calcein/Mn<sup>2+</sup> complex was added to 25  $\mu$ l of the LAMP reaction mixture. When the reaction was complete, the result was interpreted as positive if the color changed from orange to green, and negative if it did not. The results of the chromogenic reaction did not differ from the results of the real-time monitoring of turbidity.

Sensitivity of tcdA LAMP reaction

To measure the sensitivity of the primers in the LAMP detection of the *tcdA* gene, both real-time turbidity measurements and visual detection of LAMP were compared to traditional PCR. Pure genomic DNA was

Commented [LMG13]: How?

extracted from  $C.\_difficile$  VPI10463 using a Wizard Genomic DNA Purification kit, and then serially diluted 10-fold from 207 ng/µl to 0.000207 pg/µl. Distilled water was used as a negative control. As demonstrated in Fig. 4A and Fig. 4B, the detection limit of real-time turbidity was 20.7 pg/µl, which was identical to that of visual detection. The NKI and NK2 primers with the same concentration of C.difficile VPI10463 DNA were also evaluated using PCR. The detection limit for PCR was 207 pg/µl (Fig. 4C), which was 10-fold lower than that of the LAMP reaction.

#### Evaluation of the assay with stool specimens

A total of 164 stool specimens were considered eligible and suitable for this study. The subjects all showed As a common clinical presentation of CDI, pseudomembranous colitis, which is a common symptom of CDI was investigated in our inpatients and the results are shown in (Fig. 5A-B). Electronic colonoscopy revealed numerous scattered pale yellow pseudomembranes and areas of congested, brittle mucosa (Fig. 5A). Histopathological imaging (Fig. 5B) revealed an infiltration of inflammatory cells into the mucosal lamina propria. The results of LAMP and PCR detection of the todA gene in the stool specimens are compared in Table 3.

The detection rate of the tcdA gene using the LAMP method was 17% (28/164), significantly higher than the 10% (16/164) detection rate of the PCR method ( $X^2$ =47, df=1, P<0.01). The consistency between the LAMP and PCR methods was moderate (Kappa=0.533, p<0.01). Notably, 15 of the stool specimens that were negative in PCR but positive in LAMP were found to be positive in tcdA gene sequencing. Of the three specimens that were negative in LAMP but positive in PCR, only one was found to be positive using tcdA gene sequencing-results, while the other two specimens were negative. Using sequencing as the reference standard, the LAMP assay outperformed the PCR assay with a sensitivity of 97% compared to 48%, and a specificity of 100%, nearly identical to PCR's 99%. The LAMP assay achieved a PPV of 100% and an NPV of 99%, whereas the PCR assay had a PPV of 88% and an NPV of 90% (Table 4 and Table 5).

### Discussion

This study marks a significant advancement in the rapid detection of *C. difficile*, with the The optimized LAMP assay reducing the time to detect the *tcdA* gene to under 60 minutes. Incorporating loop primers (LB and LF), which are complementary to the dumbbell-shaped DNA structures, substantially reduced LAMP reaction times by over 50%, underscoring their utility in expediting the amplification process. Including loop primers also enhanced the LAMP reaction's efficiency and sensitivity by multiplying the initiation sites for amplification. Moreover, the reaction's selectivity was markedly improved as these loop primers are designed to be activated singularly during the synthesis of the artificial template (Nagamine et al., 2002). LAMP-based detection yielded negative results for all-26 pathogenic bacterial strains lacking the *tcdA* genefrom different genera than *C. difficile*, demonstrating the primers' exceptional specificity. The 10-fold higher sensitivity of the LAMP method compared to conventional PCR corroborates findings from prior studies (Kim et al., 2022; Carvajal-Gamez et al., 2023) reinforcing the method's superiority and potential to supplant conventional PCR in clinical diagnostics. These results advocate for the and supports integration of the this LAMP assay into routine diagnostic workflows, promising significant improvements in the speed and accuracy of pathogen detection.

This study demonstrated that tThe LAMP method yielded a detection rate of 17% for the *tcdA* gene of *C*. *difficile* in stool specimens, a figure that which is similar to the rates observed in diarrheal stool samples of

**Commented [LMG14]:** Did all the patients show the symptoms

**Commented [LMG15]:** How was tcdA gene sequencing done on the stool samples?

**Commented [LMG16]:** Again, how is tcdA sequencing done?

diverse populations across China. For instance, a multicenter study in Shanghai, China, reported a detection rate of 18% (93/531) for the tcdA gene (Mi et al., 2020), while an independent cross-sectional study in Southwest China noted a detection rate of 14% (125/978; Liao et al., 2018). These findings suggest that the LAMP method's efficacy in detecting the tcdA gene is comparable to the methods used in previous studies. Moreover, this study also revealed a significant discrepancy in the detection rates of the tcdA gene in stool specimens between LAMP and PCR, with LAMP exhibiting showed a superior detection rate. This was further evidenced by the fact that 15 of the stool specimens that were negative in PCR but positive in LAMP were subsequently confirmed to be true positives through sequencing of the tcdA gene. In our previous study of 300 cultured C. difficile strains from Southern Medical University Nanfang Hospital, a perfect concordance (kappa = 1) was observed between LAMP and PCR detections of the tcdA gene (Lin et al., 2022), supporting the notion that LAMP is equally reliable when bacterial DNA is present in higher quantities. The difference in consistency between LAMP and PCR in detecting C. difficile strains in stool specimens in this study may be attributed to the variable concentration of the C. difficile DNA present. In stool specimens where C. difficile DNA may be at relatively low concentrations, LAMP's lower detection limit allows for a higher detection rate. Conversely, the increased DNA content in cultured C. difficile strains following enrichment culture may mask the differences in detection limits between the two methods, leading to a high degree of consistency in detecting the tcdA gene in these strains. These findings collectively advocate for the integration of LAMP in clinical diagnostics of CDI, particularly for cases where PCR may not provide sufficient sensitivity. Building on the empirical evidence, the further statistical analysis performed in this study strengthens the case for the clinical application of LAMP. In the analysis of 164-stool samples, the LAMP assay, with its showed high sensitivity (97%) and perfect specificity (100%), along with a PPV of 100% and an NPV of 99%, demonstrated superior diagnostic accuracy when using sequencing as the gold standard. The data This suggests that the LAMP assay offers a high degree of reliability in both detecting the presence of the target pathogen and verifying its absence. In contrast, the PCR assay, despite its high specificity (99%), demonstrated significantly reduced sensitivity (48%), potentially limiting its effectiveness as a standalone diagnostic tool. The PCR's PPV of 88% and NPV of 90% suggest that while positive results are likely accurate, the risk of false negatives is increased, which could contribute to under-diagnosis. Consequently, the LAMP assay could represent a more robust approach for precise detection of CDI within the examined population. The LAMP method exhibits considerable advantages in the detection of the tcdA gene for C. difficile. First, eliminating the DNA denaturation step simplifies the operational workflow, as reactions are carried out at an isothermal condition of 60-65 °C using the strand displacement activity of Bst DNA polymerase, thus obviating the need for the precise thermal cycling equipment required for PCR (Soroka et al., 2021). This significantly lowers the technical requirements of detection, allowing molecular diagnostic techniques to be performed in resource-limited settings. Second, the LAMP reactions can be directly visualized through color changes in the calcium-magnesium complex when exposed to UV light, circumventing the need for the complex gel electrophoresis step that is requisite in PCR. This advancement not only expedites detection speed, but also minimizes the dependency on carcinogenic dyes, thereby enhancing the safety profile of the procedure. Third, the sensitivity of LAMP is 10-100 times higher than that of conventional PCR (Kim et al., 2022; Ashmi et al., 2023; Carvajal-Gamez et al., 2023). This significant enhancement in sensitivity ensures accurate detection of pathogens even at low concentrations, which is vital for early intervention and for controlling the spread of infection. Finally, the specificity of LAMP is notably enhanced through the employment of 4-6 primers, which identify up to

299

300

301

302

303

304

305

306

307

308

309

310

311

312

313

314

315

316

317

318

319

320

321

322

323

324

325

326

327

328

329

330

331

332

333

334

335

336

337

338

339

340

341

342

eight specific locations on the DNA template. This significantly mitigates the likelihood of false positives, a potential risk associated with the two-primer system in PCR (Soroka et al., 2021). In conclusion, the LAMP technique, characterized by its increased specificity and heightened sensitivity, along with a streamlined operational procedure and safe visual detection mechanism, stands out as an optimal choice for the swift and accurate detection of the *tcdA* gene. In regions constrained by economic resources and advanced diagnostic infrastructure, the benefits of the LAMP technique become even more salient. It holds potential to catalyze the broader implementation of CDI diagnostic technology, thereby contributing to an improvement in global health outcomes.

While the LAMP method stands out for its advantages, it is not without its limitations. Crosscontamination arising from multiple pipetting steps is a notable concern, particularly with material present in the aerosol. This study mitigated this issue by employing low-melting-point paraffin to seal the reaction mixtures, thereby preventing the spread of amplification products. Rigorous sample handling protocols and enhanced ventilation should be conducted to minimize contamination risks. Another limitation is the suitability of LAMP products for downstream applications, such as sequencing or cloning (Sahoo et al., 2016). This drawback underscores the need for further methodological refinements to expand the utility of LAMP products beyond mere detection. Additionally, the sensitivity of LAMP method's ability to detect inhibitors within samples remains a complex challenge (Dong et al., 2014), as it involves simultaneous inhibitor detection and material amplification. This highlights a potential area for the development of more sophisticated controls and detection metrics within the LAMP protocol. Finally, the propensity for primer-dimer formation in LAMP could lead to false positives. This underscores the importance of careful primer design and validation to ensure the reliability of the LAMP assay. Despite its transformative potential for resource limited settings, the LAMP method requires careful consideration of its limitations. Addressing these challenges is essential for harnessing the full potential of LAMP in clinical diagnostics and ensuring its role in advancing global health diagnostics.

A limitation of this study is the exclusive use of the *C. difficile* VPI10463 strain as a positive control, which may not represent the diversity of *C. difficile* strains. To address this limitation and to understand its potential impact on the conclusions drawn from the results, a more comprehensive analysis was performed. The subsequent sequencing analysis of 15 stool specimens that tested positive with the LAMP assay but negative with PCR provided additional insights. Not only did this analysis confirm the presence of *C. difficile* VPI10463 in two cases, but it also identified other strains in the remaining samples: seven cases of GZ14, two cases of ZR48, and one case each of ZR80, ZR50, GZ5, and SH8. These findings suggest that the LAMP assay developed in this study may have the capacity to detect a broader spectrum of *C. difficile* strains than initially anticipated, which is a significant consideration for the validity of this method. This understanding of the assay's limitations and their possible impact on the conclusions drawn in this study is critical for interpreting the study's results and for guiding future research directions.

#### Conclusions

In this study, the LAMP method proved to be an effective technique for the rapid and visual detection of the *tcdA* gene of *C. difficile*, demonstrating potential advantages over PCR in terms of speed, simplicity, and sensitivity. The *tcdA*-LAMP assay is particularly suitable for medical diagnostic environments with limited resources and represents a promising diagnostic strategy for the screening and detection of CDI in populations at high risk. Future research will need to further validate the applicability of the LAMP technique across various experimental and clinical settings and address any technical challenges that may arise during its broader implementation.

388 Acknowledgements 389 We would like to thank the Institute of Disease Control and Prevention and the Academy of Military 390 Medical Sciences for their technical assistance. 391 392 Informed Consent Statement 393 Because this study was based on the clinical examination of existing stool specimens for clinical research, 394 it did not require patients to provide additional samples. In the clinical analysis, all patient identification 395 information was expressed in code or pinyin to maintain the personal privacy of the patients and prevent 396 individual health information from being exposed. The test results were only used for clinical research, 397 and no test report was issued that would affect the diagnosis and treatment of the subjects. No patient risk 398 was involved in this study. Therefore, an informed consent waiver was obtained from Nanfang Hospital 399 of Southern Medical University for this study. 400 401 402 References 403 Alfa, M.J., Kabani, A., Lyerly, D., Moncrief, S., Neville, L.M., Al-Barrak, A., Harding, G.K., Dyck, B., 404 Olekson, K., and Embil, J.M. 2000. Characterization of a toxin A-negative, toxin B-positive strain of 405 Clostridium difficile responsible for a nosocomial outbreak of Clostridioides difficile-associated diarrhea. 406 JOURNAL OF CLINICAL MICROBIOLOGY 38:2706-2714. 10.1128/JCM.38.7.2706-2714.2000 407 Ashmi, M., Kumar, B., Sanjana, Abhishek, Kumar, D., and Singh, P. 2023. Rapid and specific detection 408 of B. melitensis targeting BME11661 gene using loop-mediated isothermal amplification (LAMP) 409 combined with lateral flow immunoassay (LFIA). CURRENT MICROBIOLOGY 80:351. 410 10.1007/s00284-023-03463-1 411 Badparva, E., Javadi, M.A., Kheirandish, F., Ebrahimzadeh, F., and Fallahi, S. 2022. Development and 412 evaluation of a loop-mediated isothermal amplification (LAMP) technique for rapid, accurate, and 413 specific detection of Blastocystis spp. in AIDS patients. INFECTION 50:1295-1302. 10.1007/s15010-414 022-01818-7 415 Bartlett, J.G. 2010. Detection of Clostridium difficile infection. INFECTION CONTROL AND 416 HOSPITAL EPIDEMIOLOGY 31 Suppl 1:S35-S37. 10.1086/655999 417 Brennhofer, S.A., Rogawski, M.E., Liu, J., Guerrant, R.L., Platts-Mills, J.A., and Warren, C.A. 2022. 418 Clostridioides difficile colonization among very young children in resource-limited settings. CLINICAL 419 MICROBIOLOGY AND INFECTION 28:996-1002. 10.1016/j.cmi.2022.01.022 420 Buderer, N.M. 1996. Statistical methodology: I. Incorporating the prevalence of disease into the sample 421 size calculation for sensitivity and specificity. ACADEMIC EMERGENCY MEDICINE 3:895-900. 422 10.1111/j.1553-2712.1996.tb03538.x 423 Carvajal-Gamez, B.I., Olguín-Barrera, A., Tinoco-Gracia, L., Gordillo-Perez, G., Dzul-Rosado, K., 424 Aguilar-Tipacamú, G., Hidalgo-Ruiz, M., and Mosqueda, J. 2023. Development and validation of a novel 425 detection method for Rickettsia rickettsii using a loop-mediated isothermal amplification assay. 426 FRONTIERS IN MICROBIOLOGY 14:1276809. 10.3389/fmicb.2023.1276809 427 Chen, Y.X., Lou, Y.R., Duan, L.J., Zhou, Q.J., Xu, Z.J., Chen, F.J., Chen, H.X., Xu, G.Z., Du AF, and

387

428

Chen, J. 2023. Parallel detection of multiple zoonotic parasites using a real-time fluorogenic loop-

- 429 mediated isothermal amplification-based quadruple-sample microfluidic chip. FRONTIERS IN
- 430 MICROBIOLOGY 14:1238376. 10.3389/fmicb.2023.1238376
- 431 Crobach, M.J., Planche, T., Eckert, C., Barbut, F., Terveer, E.M., Dekkers, O.M., Wilcox, M.H., and
- 432 Kuijper, E.J. 2016. European society of clinical microbiology and infectious diseases: update of the
- 433 diagnostic guidance document for Clostridium difficile infection. CLINICAL MICROBIOLOGY AND
- 434 INFECTION 22 Suppl 4:S63-S81. 10.1016/j.cmi.2016.03.010
- 435 Dong, H.J., Cho, A.R., Hahn, T.W., and Cho, S. 2014. Development of a loop-mediated isothermal
- 436 amplification assay for rapid, sensitive detection of Campylobacter jejuni in cattle farm samples.
- 437 JOURNAL OF FOOD PROTECTION 77:1593-1598. 10.4315/0362-028X.JFP-14-056
- 438 Drudy, D., Fanning, S., and Kyne, L. 2007. Toxin A-negative, toxin B-positive Clostridium difficile.
- 439 INTERNATIONAL JOURNAL OF INFECTIOUS DISEASES 11:5-10. 10.1016/j.ijid.2006.04.003
- 440 Drudy, D., Harnedy, N., Fanning, S., Hannan, M., and Kyne, L. 2007. Emergence and control of
- 441 fluoroquinolone-resistant, toxin A-negative, toxin B-positive Clostridium difficile. INFECTION
- 442 CONTROL AND HOSPITAL EPIDEMIOLOGY 28:932-940. 10.1086/519181
- 443 Freeman, J., Bauer, M.P., Baines, S.D., Corver, J., Fawley, W.N., Goorhuis, B., Kuijper, E.J., and
- 444 Wilcox, M.H. 2010. The changing epidemiology of Clostridium difficile infections. CLINICAL
- 445 MICROBIOLOGY REVIEWS 23:529-549. 10.1128/CMR.00082-09
- 446 Guh, A.Y., Mu, Y., Winston, L.G., Johnston, H., Olson, D., Farley, M.M., Wilson, L.E., Holzbauer, S.M.,
- 447 Phipps, E.C., Dumyati, G.K., Beldavs, Z.G., Kainer, M.A., Karlsson, M., Gerding, D.N., and McDonald,
- 448 L.C. 2020. Trends in U.S. burden of Clostridioides difficile
- 449 infection and outcomes. NEW ENGLAND JOURNAL OF MEDICINE 382:1320-1330.
- 450 10.1056/NEJMoa1910215
- 451 Hong-Min, Z., Jian, Y., Ying, L., Yuan, Y., Cui-Ping, W., Yu-Cheng, D., and Jia-Jia, C. 2023. Rapid
- detection of *Heterobasidion annosum* using a loop-mediated isothermal amplification assay.
- 453 FRONTIERS IN CELLULAR AND INFECTION MICROBIOLOGY 13:1134921.
- 454 10.3389/fcimb.2023.1134921
- 455 Kato, N., Ou, C.Y., Kato, H., Bartley, S.L., Brown, V.K., Dowell, V.J., and Ueno, K. 1991. Identification
- 456 of toxigenic Clostridium difficile by the polymerase chain reaction. JOURNAL OF CLINICAL
- 457 MICROBIOLOGY 29:33-37. 10.1128/jcm.29.1.33-37.1991
- 458 Kim, D.Y., Kim, H.R., Park, J.H., Kwon, N.Y., Kim, J.M., Kim, J.K., Park, J.H., Lee, K.K., Kim, S.H.,
- 459 Kim, W.I., Lyoo, Y.S., and Park, C.K. 2022. Detection of a novel porcine circovirus 4 in Korean pig
- 460 herds using a loop-mediated isothermal amplification assay. JOURNAL OF VIROLOGICAL METHODS
- 461 299:114350. 10.1016/j.jviromet.2021.114350
- 462 Kim, Y.I., Yu, C.S., Kim, Y.S., Kim, C.W., Lee, J.L., Yoon, Y.S., Park, I.J., Lim, S.B., and Kim, J.C.
- 463 2022. Clostridium difficile infection after ileostomy closure and anastomotic failure in rectal cancer
- 464 surgery patients. BJS OPEN 6. 10.1093/bjsopen/zrac026
- 465 Kuehne, S.A., Cartman, S.T., Heap, J.T., Kelly, M.L., Cockayne, A., and Minton, N.P. 2010. The role of
- 466 toxin A and toxin B in Clostridium difficile infection. NATURE 467:711-713. 10.1038/nature09397
- 467 Kuehne, S.A., Cartman, S.T., and Minton, N.P. 2011. Both, toxin A and toxin B, are important in
- 468 Clostridium difficile infection. GUT MICROBES 2:252-255. 10.4161/gmic.2.4.16109
- Kuijper, E.J., de Weerdt, J., Kato, H., Kato, N., van Dam, A.P., van der Vorm, E.R., Weel, J., van
- Kuijper, E.J., de weerdt, J., Kato, H., Kato, N., van Dam, A.P., van der vorm, E.K., weet, J., van
- 470 Rheenen, C., and Dankert, J. 2001. Nosocomial outbreak of *Clostridium difficile*-associated diarrhoea due
- 471 to a clindamycin-resistant enterotoxin A-negative strain. EUROPEAN JOURNAL OF CLINICAL
- 472 MICROBIOLOGY & INFECTIOUS DISEASES 20:528-534. 10.1007/s100960100550

- 473 Liao, F., Li, W., Gu, W., Zhang, W., Liu, X., Fu, X., Xu, W., Wu, Y., and Lu, J. 2018. A retrospective
- 474 study of community-acquired Clostridium difficile infection in southwest China. SCIENTIFIC REPORTS
- 475 8:3992. 10.1038/s41598-018-21762-7
- 476 Lin, M., Li, Z., Lin, Q., Wang, P., Liu, W., Yuan, J., Hong, Z., and Chen, Y. 2022. Development and
- 477 clinical application of a rapid and visual loop-mediated isothermal amplification test for tetM gene in
- 478 Clostridioides difficile strains cultured from feces. INTERNATIONAL JOURNAL OF INFECTIOUS
- 479 DISEASES 122:676-684. 10.1016/j.ijid.2022.07.032
- 480 Lin, M., Liu, W., Wang, P., Tan, J., Zhou, Y., Wu, P., Zhang, T., Yuan, J., and Chen, Y. 2015. Rapid
- 481 detection of ermB gene in Clostridium difficile by loop-mediated isothermal amplification. JOURNAL
- 482 OF MEDICAL MICROBIOLOGY 64:854-861. 10.1099/jmm.0.000109
- 483 Lin, Q., Pollock, N.R., Banz, A., Lantz, A., Xu, H., Gu, L., Gerding, D.N., Garey, K.W., Gonzales-Luna,
- 484 A.J., Zhao, M., Song, L., Duffy, D.C., Kelly, C.P., and Chen, X. 2020. Toxin A-predominant pathogenic
- 485 Clostridioides difficile: a novel clinical phenotype. CLINICAL INFECTIOUS DISEASES 70:2628-2633.
- 486 10.1093/cid/ciz727
- 487 Liu, W., Zou, D., Li, Y., Wang, X., He, X., Wei, X., Shao, C., Li, X., Shang, W., Yu, K., Liu, D., Li, Y.,
- 488 Guo, J., Yin, Z., and Yuan, J. 2012. Sensitive and rapid detection of the new Delhi metallo-beta-
- 489 *lactamase* gene by loop-mediated isothermal amplification. JOURNAL OF CLINICAL
- 490 MICROBIOLOGY 50:1580-1585. 10.1128/JCM.06647-11
- 491 Luo, Y., Zhang, W., Cheng, J.W., Xiao, M., Sun, G.R., Guo, C.J., Liu, M.J., Cong, P.S., and Kudinha, T.
- 492 2018. Molecular epidemiology of *Clostridium difficile* in two tertiary care hospitals in Shandong
- 493 Province, China. INFECTION AND DRUG RESISTANCE 11:489-500. 10.2147/IDR.S152724
- 494 Mi, H., Bao, R., Xiao, Y., Cui, Y., Sun, W., Shen, Y., Shi, Q., Chen, X., Lin, J., Hu, B., and Gao, X.
- 495 2020. Colonization of toxigenic Clostridium difficile among intensive care unit patients: a multi-Centre
- 496 cross-sectional study. FRONTIERS IN CELLULAR AND INFECTION MICROBIOLOGY 10:12.
- 497 10.3389/fcimb.2020.00012
- 498 Moosavian, M., Keshavarzi, R., Abbasi, M.E., and Hajiani, E. 2022. Loop mediated isothermal
- amplification of Clostridioides difficile isolates in gastrointestinal patients. AMB EXPRESS 12:42.
- 500 10.1186/s13568-022-01382-1
- 501 Mori, Y., Nagamine, K., Tomita, N., and Notomi, T. 2001. Detection of loop-mediated isothermal
- 502 amplification reaction by turbidity derived from magnesium pyrophosphate formation. BIOCHEMICAL
- 503 AND BIOPHYSICAL RESEARCH COMMUNICATIONS 289:150-154. 10.1006/bbrc.2001.5921
- Nagamine, K., Hase, T., and Notomi, T. 2002. Accelerated reaction by loop-mediated isothermal
- 505 amplification using loop primers. MOLECULAR AND CELLULAR PROBES 16:223-229.
- 506 10.1006/mcpr.2002.0415
- 507 Nawab, M., Riaz, S.K., Ismail, E., Ahamed, A., Tariq, A., Malik, M., Qusty, N.F., Bantun, F., Slama, P.,
- 508 Umair, M., Haque, S., Bonilla-Aldana, D.K., and Rodriguez-Morales, A.J. 2024. Integrated approach for
- 509 detection of SARS-CoV-2 and its variant by utilizing LAMP and ARMS-PCR. ANNALS OF CLINICAL
- 510 MICROBIOLOGY AND ANTIMICROBIALS 23:11. 10.1186/s12941-023-00665-0
- 511 Nicholson, M.R., and Donskey, C.J. 2023. Multistep testing algorithms for Clostridioides difficile
- 512 infection. JAMA 330:966-967. 10.1001/jama.2023.15875
- 513 Norén, T., Alriksson, I., Andersson, J., Akerlund, T., and Unemo, M. 2011. Rapid and sensitive loop-
- 514 mediated isothermal amplification test for Clostridioides difficile detection challenges cytotoxin B cell
- 515 test and culture as gold standard. JOURNAL OF CLINICAL MICROBIOLOGY 49:710-711.
- 516 10.1128/JCM.01824-10

- 517 Notomi, T., Okayama, H., Masubuchi, H., Yonekawa, T., Watanabe, K., Amino, N., and Hase, T. 2000.
- 518 Loop-mediated isothermal amplification of DNA. NUCLEIC ACIDS RESEARCH 28:E63.
- 519 10.1093/nar/28.12.e63
- 520 Novakova, E., Stofkova, Z., Sadlonova, V., and Hleba, L. 2021. Diagnostic methods of Clostridioides
- 521 difficile infection and Clostridioides difficile ribotypes in studied sample. ANTIBIOTICS-BASEL 10.
- 522 10.3390/antibiotics10091035
- 523 Pancholi, P., Kelly, C., Raczkowski, M., and Balada-Llasat, J.M. 2012. Detection of toxigenic
- 524 Clostridium difficile: comparison of the cell culture neutralization, Xpert C. difficile, Xpert C. difficile
- 525 /Epi, and Illumigene *C. difficile* assays. JOURNAL OF CLINICAL MICROBIOLOGY 50:1331-1335.
- 526 10.1128/JCM.06597-11
- 527 Parida, M., Sannarangaiah, S., Dash, P.K., Rao, P.V., and Morita, K. 2008. Loop mediated isothermal
- 528 amplification (LAMP): a new generation of innovative gene amplification technique; perspectives in
- 529 clinical diagnosis of infectious diseases. REVIEWS IN MEDICAL VIROLOGY 18:407-421.
- 530 10.1002/rmv.593
- 531 Rodríguez, C., Romero, E., Garrido-Sanchez, L., Alcaín-Martínez, G., Andrade, R.J., Taminiau, B.,
- 532 Daube, G., and García-Fuentes, E. 2020. Microbiota insights in Clostridium difficile infection and
- 533 inflammatory bowel disease. GUT MICROBES 12:1725220. 10.1080/19490976.2020.1725220
- 534 Rupnik, M. 2008. Heterogeneity of large clostridial toxins: importance of Clostridium difficile
- 535 toxinotypes. FEMS MICROBIOLOGY REVIEWS 32:541-555. 10.1111/j.1574-6976.2008.00110.x
- 536 Sahoo, P.R., Sethy, K., Mohapatra, S., and Panda, D. 2016. Loop mediated isothermal amplification: An
- innovative gene amplification technique for animal diseases. VETERINARY WORLD 9:465-469.
- 538 10.14202/vetworld.2016.465-469
- 539 Shuai, H., Bian, Q., Luo, Y., Zhou, X., Song, X., Ye, J., Huang, Q., Peng, Z., Wu, J., Jiang, J., and Jin, D.
- 540 2020. Molecular characteristics of *Clostridium difficile* in children with acute gastroenteritis from
- 541 Zhejiang. BMC INFECTIOUS DISEASES 20:343. 10.1186/s12879-020-05030-6
- 542 Soroka, M., Wasowicz, B., and Rymaszewska, A. 2021. Loop-mediated isothermal amplification
- 543 (LAMP): the better sibling of PCR?. CELLS 10. 10.3390/cells10081931
- 544 Tang, C., Cui, L., Xu, Y., Xie, L., Sun, P., Liu, C., Xia, W., and Liu, G. 2016. The incidence and drug
- 545 resistance of *Clostridium difficile* infection in Mainland China: a systematic review and meta-analysis.
- 546 SCIENTIFIC REPORTS 6:37865. 10.1038/srep37865
- 547 Ushikubo, H. 2004. Principle of LAMP method--a simple and rapid gene amplification method. UIRUSU
- 548 54:107-112. 10.2222/jsv.54.107

556

- 549 Wen, B.J., Dong, N., Ouyang, Z.R., Qin, P., Yang, J., Wang, W.G., Qiang, C.X., Li, Z.R., Niu, Y.N., and
- 550 Zhao, J.H. 2023. Prevalence and molecular characterization of Clostridioides difficile infection in China
- 551 over the past 5 years: a systematic review and meta-analysis. INTERNATIONAL JOURNAL OF
- 552 INFECTIOUS DISEASES 130:86-93. 10.1016/j.ijid.2023.03.009
- 553 Yu, L., Li, H., Zhao, X., Wang, X., Wei, X., Lin, W., Li, P., Cui, L., and Yuan, J. 2017. Rapid visual
- 554 detection of binary toxin producing *Clostridium difficile* by loop-mediated isothermal amplification.
- 555 EXPERIMENTAL AND THERAPEUTIC MEDICINE 14:4781-4788. 10.3892/etm.2017.5178