Check for updates

#### **OPEN ACCESS**

EDITED BY Tim Dumonceaux, Agriculture and Agri-Food Canada (AAFC), Canada

REVIEWED BY Amir Sohrabi, Karolinska Institutet (KI), Sweden M. Jahangir Alam, University of Houston, United States

\*CORRESPONDENCE Pengbo Guo 🖾 877273621@qq.com Yinghui Zhao 🖾 zhaoyinghui1999@sina.com

<sup>†</sup>These authors have contributed equally to this work

SPECIALTY SECTION This article was submitted to Microbiotechnology, a section of the journal Frontiers in Microbiology

RECEIVED 10 November 2022 ACCEPTED 27 February 2023 PUBLISHED 23 March 2023

#### CITATION

Liu W, Lu G, Wang Y, Chen Z, Gao Y, Yin Z, Wu Y, Lv X, Guo P and Zhao Y (2023) A novel loop-mediated isothermal amplification-lateral flow dipstick method for *Helicobacter pylori* detection. *Front. Microbiol.* 14:1094600. doi: 10.3389/fmicb.2023.1094600

#### COPYRIGHT

© 2023 Liu, Lu, Wang, Chen, Gao, Yin, Wu, Lv, Guo and Zhao. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

# A novel loop-mediated isothermal amplification-lateral flow dipstick method for *Helicobacter pylori* detection

Wenwen Liu<sup>1†</sup>, Gang Lu<sup>2,3†</sup>, Yu Wang<sup>2,3</sup>, Zhenghong Chen<sup>4</sup>, Yunyun Gao<sup>2</sup>, Zhipeng Yin<sup>2</sup>, Yi Wu<sup>2</sup>, Xiaoqian Lv<sup>2</sup>, Pengbo Guo<sup>2,3\*</sup> and Yinghui Zhao<sup>2,3\*</sup>

<sup>1</sup>Department of Clinical Laboratory Medicine, The First Affiliated Hospital of Shandong First Medical University and Shandong Provincial Qianfoshan Hospital, Shandong Medicine and Health Key Laboratory of Laboratory Medicine, Jinan, Shandong, China, <sup>2</sup>Department of Pathogen Biology, School of Clinical and Basic Medical Sciences, Shandong First Medical University and Shandong Academy of Medical Sciences, Jinan, Shandong, China, <sup>3</sup>Department of Clinical Microbiology Laboratory, Shandong Provincial Hospital Affiliated to Shandong First Medical University, Jinan, Shandong, China, <sup>4</sup>Key Laboratory of Medical Microbiology and Parasitology, Department of Microbiology, Guizhou Medical University, Guiyang, China

**Introduction:** To eradicate *Helicobacter pylori* (*H. pylori*) and reduce the risk of gastric cancer, a sensitive, specific, convenient, and simple detection method is needed. This study aimed to establish a novel loop-mediated isothermal amplification-lateral flow dipstick (LAMP-LFD) method for *H. pylori* detection.

**Methods:** LAMP primer design software was used to design primers for the conserved sites of the *H. pylori* ureB gene. UreB-FIP-labeled biotin was used for LAMP amplification, and FAM-labeled probes were specifically hybridized with LAMP amplification products, which were then detected by LFD. In addition, a clinical study was conducted to assess LAMP-LFD in 20 fecal samples.

**Results:** The results of the optimization indicated that *H. pylori* could be specifically detected by LFD without cross-reaction with other non-*H. pylori* bacteria when the LAMP was performed at 65°C for 60 min. The lower limit of the detection method was  $10^2$  copies/µL, which was 100 times the sensitivity of polymerase chain reaction (PCR). *H. pylori*-positive fecal samples were detected by LAMP-LFD in 13/20 patients.

**Discussion:** In conclusion, a new LAMP-LFD assay has been fully established and confirmed for *H. pylori* detection. The entire process can be completed in approximately 1.5 h, with the advantages of strong specificity, high sensitivity, and simple operation. This study provides a novel potential method for the detection of *H. pylori* in the clinical settings of primary hospitals and low-resource countries.

KEYWORDS

Helicobacter pylori, ureB gene, loop-mediated isothermal amplification, lateral flow dipstick, novel detection method

# Introduction

*Helicobacter pylori* (*H. pylori*), defined as a class I carcinogen by the World Health Organization, is the most common pathogen that infects the stomach and often resides on the surface of the pylorus and duodenal mucosa (Hooi et al., 2017). More than 50% of the world's population is infected with it, and the infection rate in developing countries is generally higher than that in developed countries (Roszczenko-Jasińska et al., 2020). It can cause various gastrointestinal and extragastric diseases, such as gastritis, gastric ulcers, gastric cancer, mucosa-associated lymphoid tissue lymphoma, Alzheimer's disease, and other

10.3389/fmicb.2023.1094600

neurological or blood diseases (Kountouras et al., 2010; Kutlubay et al., 2014; Sacc et al., 2014; Sarem and Corti, 2016; Muhammad et al., 2017; Zamani et al., 2017; Mladenova, 2019; Pellicano et al., 2020; Howden, 2021). Early eradication of the bacterium is important to prevent and control these diseases. Therefore, it is crucial to develop a rapid, accurate, and suitable method for *H. pylori* detection.

Techniques for *H. pylori* detection can be classified into invasive and non-invasive approaches (Mohammadian and Ganji, 2019). Rapid urease test (RUT), histological examination, and bacterial isolation or culture are considered invasive methods, while PCR is considered a non-invasive method (Di Bonaventura et al., 2004; Patel et al., 2014; Milani et al., 2019; Abdelmalek et al., 2022). Each approach has its advantages and limitations. Of the non-invasive methods, PCR has good sensitivity and specificity; however, it requires PCR and gel imaging equipment, limiting its use in primary hospitals or low-resource countries. To overcome these difficulties, a new method must be developed.

Loop-mediated isothermal amplification (LAMP) is a novel gene amplification strategy for continuous extension and replacement of four primer pairs within 1h under certain temperature conditions and strand displacement by Bst DNA polymerase (Liu et al., 2019). This method has many advantages, including simplicity, speed, high sensitivity and specificity, and minimal equipment or reagent requirements (Domesle et al., 2018). The amplification products of LAMP are mainly detected by turbidimetry, electrophoresis, or calcein fluorescent dyes (Bakhtiari et al., 2016, 2019; Park et al., 2018; Sohrabi et al., 2021). The contact of ethidium bromide (EB), a carcinogen, the potential harm from calcein fluorescent dye to the human body, and the subjectivity of the turbidimeter reduce the practicality of LAMP in the field (Zhang et al., 2019). Lateral flow dipstick (LFD) can overcome these limitations (Lalle et al., 2018). Consequently, this study combined LAMP with LFD to establish a LAMP-LFD method for the detection of *H. pylori*, which will provide a reliable and effective diagnostic tool for H. pylori infection.

# Materials and methods

### **Experimental materials**

A bacterial DNA extraction kit was obtained from Dalian Bao Biotechnology Co., Ltd. A PCR amplification kit and a Bst DNA polymerase were obtained from Kangwei Century Biotechnology Co., Ltd. Hydroxynaphthalene blue (HNB) was obtained from Sigma Co., Ltd. The dNTP mixture was obtained from SOLEBAO Biotechnology Co., Ltd. A <sup>14</sup>C Urea breath test (UBT) kit was obtained from Shenzhen Zhonghe Haidewei Biotechnology Co., Ltd. A 5kb DNA marker was obtained from Nuovizan Biotechnology Co., Ltd. *H. pylori* (ATCC26695), *Escherichia coli* (DH5 $\alpha$ ), and *Pseudomonas aeruginosa* (BNCC186335) were preserved in our laboratory. *Staphylococcus aureus* was presented by Jing Ni, a teacher from the School of Public Health of Shandong First Medical University. The fecal samples of patients with gastric diseases were collected from the Oncology Department of Taishan Hospital. The fecal samples experiments were approved by the Ethics Committee of Shandong First Medical University (R202211250164).

#### Bacterial culture and DNA extraction

*H. pylori* (ATCC26695) was cultured on a yolk agar medium under the conditions of 85% N<sub>2</sub>, 10% CO<sub>2</sub>, and 5% O<sub>2</sub> at  $37^{\circ}$ C for 3 d. Other bacteria (*Escherichia coli, Pseudomonas aeruginosa*, and *Staphylococcus aureus*) were cultured on ordinary agar medium at  $37^{\circ}$ C for 20 h under the same gas conditions. Bacterial DNA was extracted according to the kit (TaKaRa MiniBEST Bacteria Genomic DNA Extraction Kit Ver.3.0, TaKaRa) instructions. The concentration and purity of the DNA were evaluated using a NanoDrop UV spectrophotometer with A<sub>260</sub> and A<sub>280</sub> absorbance values.

### Primer and probes

The H. pylori ureB gene sequence was obtained from NCBI (sequence number: WP\_ 000724295.1, AFV41291.1, and AY714224.1). The conserved region of the ureB gene was selected by Mega and Bio edit software. Primer Explorer V5 was used to design several groups of LAMP primers and probes, and a group with high specificity was selected. Each group of primers and probes included two internal primers (FIP, 5'-CCAGCACCTTCAGTG TGGAAATTGCGTGGAAGACACTATGG-3' and BIP 5' -GGCG GACACGCTCCTGATATTTGGAAGCGGGAAGAATGTTG-3'), two external primers (F3, 5'-GTCGCTATCCACACAGACAC-3' and B3, 5'-AGTGAAAGGGATAGTGGGGT-3'), and a probe (Hp, 5'-CCGGACGCACTATGCACAC-3') in which the 5' end of FIP was labeled with biotin, while the 5' end of Hp was labeled with FAM. The primers and probes were synthesized and labeled by Bioengineering Co., Ltd. (Shanghai, China). PCR primers (UreB-F, 5'-GAACATGACTACACCAT-3' and UreB-R, 5'-TGGTTTGAGGGCGAATC-3') were synthesized by the aforementioned company.

# Establishment and optimization of the LAMP reaction system

The 25  $\mu$ L reaction system consisted of 2.5  $\mu$ L 10× reaction buffer, 6 mM MgSO<sub>4</sub>, 1.4 mM dNTPs, 1.6  $\mu$ M FIP, 1.6  $\mu$ M BIP, 0.2  $\mu$ M F3, 0.2  $\mu$ M B3, 0.32 U/ $\mu$ L Bst DNA polymerase, 1  $\mu$ L *H. pylori* DNA template, and 11.5  $\mu$ L ddH<sub>2</sub>O. The solutions were added to a 200  $\mu$ L reaction tube. The tubes were placed in a 65°C water bath for 1 h, followed by a 85°C water bath for 20 min. Reaction products were verified by 2.5% agarose gel electrophoresis.

Next, the LAMP reaction conditions were optimized. The dNTP mixture concentration was tested from 4 mM to 14 mM, with the concentration increased at 2 mM intervals. The LAMP amplification products were analyzed using gel electrophoresis. The dNTP concentration with the most obvious electrophoretic band was selected to optimize the  $Mg^{2+}$  content in the reaction. The

 $Mg^{2+}$  content was tested from 0 to 3.0 µL at 0.5 µL intervals. LAMP amplification was performed, and the products were detected using gel electrophoresis. Bst DNA polymerase content was optimized according to the dNTP and Mg<sup>2+</sup> optimized reaction system. Bst DNA polymerase content was tested from 0.5 to 2.0  $\mu$ L at 0.5  $\mu$ L intervals. LAMP amplification was performed, and the products were observed after gel electrophoresis. The concentration ratio of the internal and external primers was then optimized. The concentration of external primers (F3, B3) was maintained at 5 µM, while the concentrations of internal primers (FIP and BIP) were diluted to 50, 40, 20, 10, and  $5\,\mu$ M; the concentration ratios of internal and external primers were 10:1, 8:1, 4:1, 2:1, and 1:1, respectively. After the reaction, agarose gel electrophoresis was performed for verification purposes. Based on the earlier optimized reaction system, the reaction time and temperature were optimized. The reaction temperatures were tested at 50, 55, 60, 65, and 70°C. The reaction time was set from 15 to 75 min at 15-min intervals. The optimal reaction temperature and time were selected to correspond to the most obvious electrophoresis band using 2.5% gel electrophoresis analysis.

### Establishment of LAMP-LFD

The optimized LAMP reaction system was then used with the biotin-labeled FIP and FAM-labeled probe. The reaction system included a 1 µL FAM-labeled probe (20 pmol/L), hybridized at 65°C for 5 min, and then heated at 85°C for 20 min. The reaction products (9 µL) were added to 190 µL ddH<sub>2</sub>O, and the diluted reaction product was added to the sample area of the LFD. The product migrated forward to the sample-binding pad, detection line, and quality control line. The DNA probe labeled by biotin-FAM effectively bound to streptavidinized colloidal gold on the sample-binding pad. If biotin-FAM probe amplification of the target gene occurred, the product would be captured by the anti-FAM antibody to form a ternary complex, which would show a red band on the detection line. In contrast, biotin-FAM-labeled probes that do not form hybrid products would cross the detection line and combine with biotinylated BSA on the quality control line, showing only red bands on the quality control line.

# Analysis of specificity, sensitivity, and repeatability of LAMP-LFD

To verify the specificity of LAMP-LFD, DNA from *H. pylori*, *Escherichia coli*, *Pseudomonas aeruginosa*, and *Staphylococcus aureus* was extracted. The optimal LAMP assay was used for amplification. Amplification products were detected using LFD and 2.5% agarose gel electrophoresis.

To test the sensitivity of the method, the concentration of *H. pylori* DNA was diluted with sterile ddH<sub>2</sub>O with a 10-fold gradient for a final concentration range of  $10-10^8$  copies/µL. Different concentrations of DNA were used as templates to perform the optimized LAMP amplification reaction, and the amplification products were detected using LFD and 2.5% agarose gel electrophoresis. At the same time, PCR was conducted using

the following reaction system:  $25 \ \mu L \ 2 \times Es$  Taq Master Mix (Dye), 2  $\mu$ L ureB-F primer, 2  $\mu$ L ureB-R primer, 1  $\mu$ L template DNA, and 20  $\mu$ L ddH<sub>2</sub>O. Cycling conditions were as follows: denaturation at 94°C for 2 min, 30 cycles of denaturation at 94°C for 30 s, annealing at 56°C for 30 s, elongation at 72°C for 30 s, and final elongation at 72°C for 2 min. The PCR products were detected using 1.5% agarose gel electrophoresis.

Three groups of parallel samples were prepared to test the repeatability of this method. *H. pylori* DNA was diluted to  $10^4$  copies/µL using ddH<sub>2</sub>O. The optimized LAMP reaction system was prepared using diluted templates. Finally, LFD and 2.5% agarose gel electrophoresis were used to detect the amplification products in the three groups of samples.

# The detection of clinically isolated strains by LAMP-LFD

Fecal samples from 20 patients with gastric disease were collected. *H. pylori* DNA from feces was extracted using the kit. LAMP-LFD was used to detect bacterial DNA. PCR and <sup>14</sup>C-urea breath test (<sup>14</sup>C-UBT) were conducted simultaneously.

# Results

#### Optimization of the LAMP reaction system

The concentration of dNTPs was optimized first. The target sequence was amplified when the dNTP concentration was 4-14 mM (Figure 1A); bands were unclear when the concentration was lower than 4 mM. When the dNTP concentration was 8 mM, the electrophoretic band exhibited an obvious ladder shape. The electrophoretic band did not change significantly when the concentration was between 8 and 10 mM. Consequently, 8 mM was selected as the optimal concentration of dNTPs. Optimization of the Mg<sup>2+</sup> concentration was then performed. There were no bands in the 0 and 0.5  $\mu L~Mg^{2+}$  groups (Figure 1B). When the  $Mg^{2+}$ concentration was 1–2.5  $\mu$ L, the target bands were observed, and the band of the 1.5  $\mu$ L group was the most intense. The target band was not detected when the  $Mg^{2+}$  concentration was 3.0  $\mu$ L; the specificity of the reaction might have been reduced due to the high Mg<sup>2+</sup> content. Therefore, it was determined that the most suitable  $Mg^{2+}$  concentration in the reaction system was 1.5  $\mu L$ 

Enzyme concentration optimization experiments showed that the electrophoresis band was bright and clear when the enzyme concentration was 1.0  $\mu$ L (Figure 1C). However, with increasing enzyme concentration, the brightness and clarity of the electrophoresis bands did not change. Consequently, 1  $\mu$ L of Bst DNA polymerase was found to be the optimal concentration in the reaction system.

The LAMP reaction was performed to optimize primer concentration, and the electrophoresis results revealed that bands were not obvious when the concentration ratio of the internal and external primers was 1:1 (Figure 1D). The brightness and clarity of the bands gradually improved with an increase in the ratio of the internal and external primers; the bands showed the highest intensity when the ratio was 8:1. This did not increase further



Optimization of the LAMP reaction. (A) Optimization of dNTPs concentration, M: DNA Maker DL5000; lane 1: Negative control; lane 2: 4 mM; lane 3: 6 mM; lane 4: 8 mM; lane 5: 10 mM; lane 6: 12 mM; lane 7: 14 mM. (B) Optimization of  $Mg^+$  content, M: DNA Maker DL5000; lane 1: Negative control; lane 2: 0  $\mu$ L; lane 3: 0.5  $\mu$ L; lane 6: 1.0  $\mu$ L; lane 5: 1.5  $\mu$ L; lane 6: 2.0  $\mu$ L; lane 7: 2.5  $\mu$ L; lane 8: 3.0  $\mu$ L. (C) Optimization of enzyme content, M: DNA Maker DL5000; lane 1: Negative control; lane 2: 0.5  $\mu$ L; lane 6: 2.0  $\mu$ L; lane 6: 2.0  $\mu$ L; lane 7: 2.5  $\mu$ L; lane 8: 3.0  $\mu$ L. (C) Optimization of enzyme content, M: DNA Marker DL5000; lane 1: Negative control; lane 2: 0.5  $\mu$ L; lane 6: 2.0  $\mu$ L; lane 4: 1.5  $\mu$ L; lane 5: 2.0  $\mu$ L. (D) Optimization of primer concentration ratio (internal primer: external primer), M: DNA Marker DL5000; lane 1: Negative control; lane 2: 1.1; lane 3: 2.1; lane 4: 4.1; lane 5: 1.0  $\mu$ L; lane 5: 0.0; lane 1: Negative control; lane 2: 5.5 °C; lane 5: 6.5 °C; lane 5: 6.5 °C; lane 5: 6.5 °C; lane 5: 6.5 °C; lane 6: 70 °C. (F) Optimization of reaction time, M: DNA Maker DL5000; lane 1: Negative control; lane 2: 1.5 min; lane 3: 30 min; lane 4: 45 min; lane 5: 60 min; lane 6: 75 min. The lanes marked red are the optimal conditions.

when the concentration ratio of the internal and external primers was increased to 10:1. Therefore, the internal primer and external primer concentrations were determined to be  $40 \,\mu\text{M}$  and  $5 \,\mu\text{M}$ , respectively (the ratio of the internal and external primers was 8:1).

The results of electrophoresis following temperature optimization showed that target bands were observed when the reaction temperature was 50–  $65^{\circ}$ C, whereas no target bands were found after electrophoresis when the reaction temperature rose to  $70^{\circ}$ C (Figure 1E). There was no significant difference between 50 and  $55^{\circ}$ C, and the bands were dim because the amplification efficiency of the reaction was low. When the temperature was increased to  $60^{\circ}$ C, the bands were bright. However, the  $65^{\circ}$ C group had the brightest bands, indicating that  $65^{\circ}$ C was the optimal reaction temperature.

Finally, reaction time was optimized. No target bands were observed when the reaction time was 15 min (Figure 1F). The bands were the brightest and presented an obvious ladder shape when the reaction time was increased to 60 min. Therefore, 60 min was selected as the optimal time.

# Establishment of LAMP-LFD

The positive sample showed red bands on both the detection line and the quality control line, whereas the



negative sample only showed red bands on the quality control line (Figure 2), indicating that the LAMP-LFD was successfully established.





# Specificity of LAMP-LFD

The established LAMP-LFD assay was used to detect *E. coli*, *S. aureus*, and *P. aeruginosa* (Figure 3). When *H. pylori* DNA was used as the template, the detection and quality control lines of the horizontal chromatography strip showed two obvious red bands and the electrophoresis result showed typical ladder-shaped bands, indicating a positive detection result. In contrast, no electrophoretic bands were observed when ddH<sub>2</sub>O and other bacterial DNA samples were used as templates. Simultaneously, no red bands were observed on the detection line of the horizontal chromatographic test strip, indicating that the test result was negative. The results showed that the established LAMP-LFD had good specificity.

# Sensitivity of LAMP-LFD

*H. pylori* DNA was diluted with a 10-fold gradient to obtain a DNA concentration of 10 to  $10^8$  copies/µL. LAMP was performed

under the optimized conditions, and the reaction products were detected using LFD and agarose gel electrophoresis (Figures 4B, C). The lowest detection concentration was  $10^2$  copies/µL. The same concentration of DNA was used to perform the PCR (Figure 4A). The lowest concentration detected using PCR was  $10^4$  copies/µL. The concentration of DNA detected by LAMP-LFD was 100 times higher than that detected by PCR. Therefore, the LAMP-LFD method established in this study could be used for *H. pylori* detection.

### Repeatability of LAMP-LFD

In total, three samples of *H. pylori* DNA were separately diluted to  $10^4$  copies/µL and used to perform LAMP assays. The amplified products were detected using LFD and electrophoresis. Positive results were obtained for the three parallel reactions (Figure 5), indicating that LAMP-LFD was repeatable.



### Clinical sample detection by LAMP-LFD

The established LAMP-LFD, PCR, and <sup>14</sup>C-UBT assays were used to simultaneously detect *H. pylori* in 20 clinical fecal samples. A total of 13 *H. pylori*-positive cases were detected by LAMP-LFD, 11 by PCR, and 15 by <sup>14</sup>C-UBT.

### Discussion

*H. pylori* is known to be the most important biological factor of gastric cancer, with high infection and recurrence rates worldwide (Yan and Jun, 2019). Early detection of *H. pylori* infection, fast treatment, and prevention of its transmission to other people are important strategies to reduce its harm. Currently, numerous detection methods have been established, but each diagnostic method has one or more disadvantages, such as difficult operation, time consumption, insufficient sensitivity or specificity, expensive instruments or reagents, harm to the human body, and environmental pollution (Ju and Kim, 2015; Wang et al., 2015; Syahniar et al., 2019). Therefore, it is necessary to develop new methods for overcoming these limitations.

The LAMP-LFD method described here is the first of its kind for the detection of *H. pylori* infection. The ureB gene of the bacterium was used as the target gene to establish this method. Specific LAMP primers and LFD detection probes were designed, and the reaction parameters were optimized. The LAMP assay relies on four specific primers for the identification of six gene regions in the target gene without non-specific amplification, compared with PCR using two primers to complete nucleic acid amplification (Liu et al., 2019). Previous studies have shown that the primer, Mg<sup>2+</sup>, and enzyme concentration can directly affect the specificity and sensitivity of the detection method (Mori et al., 2001). Therefore, the reaction parameters of LAMP were optimized to ensure the high specificity and sensitivity of the method.

The LAMP-LFD assay detected a minimum concentration of *H. pylori* of  $10^2$  copies/ $\mu$ L, which suggested that its sensitivity was 100 times that of PCR. LAMP-LFD amplification products

were detected using a horizontal chromatography strip with probe markers. Negative or positive results were judged according to the presence of a red strip at the detection line. The detection method could be performed without calcein and hydroxynaphthalene blue dyes, eliminating errors caused by subjective color changes and reducing contact with carcinogenic EB when preparing agarose gel electrophoresis (Domesle et al., 2018; Bakhtiari et al., 2019). In summary, the method takes advantage of the high sensitivity and specificity of LAMP, as well as the ease of display and safety of LFD. Although various molecular detection methods for H. pylori have been developed in recent years (Molnar et al., 2008; Taborda et al., 2018; Ji et al., 2020), these methods require special instruments, such as PCR, fluorescent quantitative PCR, and high-throughput sequencing, which restrict the implementation of H. pylori detection in primary hospitals or low-resource countries. Compared with other molecular methods, LAMP-LFD has the advantages of simplicity, fast operation, minimal experimental equipment requirements, less consumables, and intuitive results.

Fecal samples were analyzed by LAMP-LFD, PCR, and <sup>14</sup>C-UBT. The positive detection rate of LAMP-LFD was higher than that of PCR and lower than that of <sup>14</sup>C-UBT. The LAMP-LFD method established is a fast, simple, effective, specific, and sensitive technique for the identification of *H. pylori*. Therefore, LAMP-LFD may substitute PCR and <sup>14</sup>C-UBT for the detection of *H. pylori* infection. Furthermore, this method avoids environmental contamination and harm caused by radioactive <sup>14</sup>C in children. In addition, it is a non-invasive approach and is applicable to almost all groups, such as the elderly, children, and people with respiratory diseases.

# Conclusion

In conclusion, a LAMP-LFD method for *H. pylori* detection was successfully developed in this study. This method has outstanding accuracy, high specificity, and sensitivity and is simple, fast, and independent of expensive instruments or reagents. The entire process can be completed within 1.5 h, depending only on the water bath and the test strip. It has been developed for the selective detection of *H. pylori* in feces, which is a non-invasive detection method suitable for use in primary hospitals or low-resource countries. This study provides novel strategies for *H. pylori* detection that are worth popularizing.

## Data availability statement

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number(s) can be found in the article/ supplementary material.

# Author contributions

YZ and PG conceived and designed the study and critically revised the manuscript. WL carried out the experiments. GL drafted the manuscript. YWa, ZC, YG, ZY, YWu, and XL contributed to the revision of the manuscript. All authors read and approved the final manuscript.

# References

Abdelmalek, S., Shokry, K., Hamed, W., Abdelnaser, M., Aboubakr, A., Elenin, S. A., et al. (2022). The validity evaluation of different 16srRNA gene primers for helicobacter detection urgently requesting to design new specific primers. *Sci Rep.* 12, 10737. doi: 10.1038/s41598-022-14600-4

Bakhtiari, S., Alvandi, A., Pajavand, H., Navabi, J., Najafi, F., Abiri, R., et al. (2016). Development and diagnostic evaluation of loop-mediated isothermal amplification using a new gene target for rapid detection of *Helicobacter pylori*. *Jundishapur J Microbiol.* 9, e28831. doi: 10.5812/jjm.28831

Bakhtiari, S., Hasanvand, B., Pajavand, H., Alvandi, A., and Abiri, R. (2019). Rapid and accurate detection of *Helicobacter pylori* from biopsy specimens using loopmediated isothermal amplification. *APMIS*. 127, 510–514. doi: 10.1111/apm.12953

Di Bonaventura, G., Neri, M., Angelucci, D., Rosini, S., Piccolomini, M., Piccolomini, R., et al. (2004). Detection of *Helicobacter Pylori* by PCR on Gastric Biopsy Specimens Taken for CP Test: Comparison with Histopathological Analysis. *Int J Immunopathol Pharmacol.* 17, 77–82. doi: 10.1177/039463200401700111

Domesle, K. J., Yang, Q., Hammack, T. S., and Ge, B. (2018). Validation of a Salmonella loop-mediated isothermal amplification assay in animal food. *Int J Food Microbiol.* 264, 63–76. doi: 10.1016/j.ijfoodmicro.2017.10.020

Hooi, J. K., Lai, W. Y., Ng, W. K., Suen, M. M., Underwood, F. E., Tanyingoh, D., et al. (2017). Global prevalence of *Helicobacter pylori* infection: systematic review and meta-analysis. *Gastroenterology*. 153, 420–429. doi: 10.1053/j.gastro.2017.04.022

Howden, C. W. D. Y. (2021). Graham recent developments pertaining to h pylori infection. *Am J Gastroenterol*. 116, 1–3. doi: 10.14309/ajg.00000000001031

Ji, Y. J., Liang, X., and Lu, H. (2020). Analysis of by high-throughput sequencing: *Helicobacter pylori* infection and salivary microbiome. *BMC Oral Health.* 20, 84. doi: 10.1186/s12903-020-01070-1

Ju, Y. L., and Kim, N. (2015). Diagnosis of *Helicobacter pylori* by invasive test: histology. *Ann Transl Med.* 3, 10. doi: 10.3978/j.issn.2305-5839.2014.11.03

Kountouras, J., Boziki, M., Gavalas, E., Zavos, C., Deretzi, G., Chatzigeorgiou, S., et al. (2010). Five-year survival after *Helicobacter pylori* eradication in Alzheimer disease patients. *Cognit. Behav. Neurol.* 23, 199–204. doi: 10.1097/WNN.0b013e3181df3034

Kutlubay, Z., Zara, T., Engin, B., Serdaroglu, S., Tüzün, Y., Yilmaz, E., et al. (2014). *Helicobacter pylori* infection and skin disorders. *Hong Kong Med J.* 20, 317-324. doi: 10.12809/hkmj134174

# Funding

This study was supported by the National Foundation of Natural Science of China (Grant No. 81602455), the Science Foundation of Shandong Province (Grant No. ZR2013HM033), and the Taiwan Science and Technology Innovation Development Project (Grant Nos. 2020NS136 and 2021NS406).

# **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.

### Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Lalle, M., Possenti, A., Dubey, J. P., and Pozio, E. (2018). Loop-mediated isothermal amplification-lateral-flow dipstick (LAMP-LFD) to detect *Toxoplasma gondii* oocyst in ready-to-eat salad. *Food Microbiol.* 70, 137–142. doi: 10.1016/j.fm.2017.10.001

Liu, Z. K., Zhang, Q. Y., Yang, N. N., Xu, M. G., and Xu, J. F. (2019). Rapid and Sensitive detection of salmonella in chickens using loop-mediated isothermal amplification combined with a lateral flow dipstick. *J. Microbiol. Biotechnol.* 29, 454–464. doi: 10.4014/jmb.1712.12010

Milani, M., Moaddab, Y., and Sharifi, Y. (2019). One piece biopsy for both rapid urease test and cultivation of *Helicobacter pylori*. J. Microbiol. Methods. 164, 105674. doi: 10.1016/j.mimet.2019.105674

Mladenova, I. (2019). Helicobacter pylori and cardiovascular disease: update 2019. Minerva Cardioangiol. 67, 425–432. doi: 10.23736/S0026-4725.19.04986-7

Mohammadian, T., and Ganji, L. (2019). The diagnostic tests for detection of *Helicobacter pylori* Infection. *Monoclon Antib Immunodiagn Immunother.* 38, 1–7. doi: 10.1089/mab.2018.0032

Molnar, B., Szoke, D., Ruzsovics, A., and Tulassay, Z. (2008). Significantly elevated *Helicobacter pylori* density and different genotype distribution in erosions as compared with normal gastric biopsy specimen detected by quantitative real-time PCR. *Eur J Gastroenterol Hepatol.* 20, 305–313. doi: 10.1097/MEG.0b013e3282f2fda4

Mori, Y., Nagamine, K., Tomita, N., and Notomi, T. (2001). Detection of loop-mediated isothermal amplification reaction by turbidity derived from magnesium pyrophosphate formation. *Biochem Biophys Res Commun.* 289, 150–154. doi:10.1006/bbrc.2001.5921

Muhammad, J. S., Zaidi, S. F., Saeed, S. A., and Ishaq, M. (2017). Current status of *Helicobacter pylori* association with haematological and cardiovascular diseases: a mini review. *J Pak Med Assoc.* 67, 907–911.

Park, Y. R., Kim, H. R., Kim, S. H., Lee, K. K., Lyoo, Y. S., Yeo, S. G., et al. (2018). Loop-mediated isothermal amplification assay for the rapid and visual detection of Novel Porcine Circovirus 3. J virol methods. 253, 26–30. doi:10.1016/j.jviromet.2017.12.006

Patel, S. K., Pratap, C. B., Jain, A. K., Gulati, A. K., and Nath, G. (2014). Diagnosis of *Helicobacter pylori*: what should be the gold standard? *World J Gastroenterol.* 20, 12847–12859. doi: 10.3748/wjg.v20.i36.12847

Pellicano, R., Ianiro, G., and Fagoonee, S. (2020). Review: extragastric diseases and *Helicobacter pylori. Helicobacter* 25, e12741. doi: 10.1111/hel.12741

Roszczenko-Jasińska, P., Wojty, M. I., and Jagusztyn-Krynicka, E. K. (2020). *Helicobacter pylori* treatment in the post-antibiotics era-searching for newdrug targets. *Appl Microbiol Biotechnol.* 104, 9891–9905. doi: 10.1007/s00253-020-10945-w

Sacc,à, S. C., Vagge, A., Pulliero, A., and Izzotti, A. (2014). *Helicobacter pylori* infection and eye diseases: a systematic review. *Medicine*. 93, e216. doi: 10.1097/MD.00000000000216

Sarem, M., and Corti, R. (2016). Role of *Helicobacter pylori* coccoid forms in infection and recrudescence. *Gastroenterol Hepatol.* 39, 28–35. doi: 10.1016/j.gastrohep.2015.04.009

Sohrabi, A., Franzen, J., Tertipis, N., Zagai, U., Li, W., Zheng, Z., et al. (2021). Efficacy of loop-mediated isothermal amplification for *H. pylori* detection as point-of-care testing by noninvasive sampling. *Diagnostics*. 11, 1538. doi: 10.3390/diagnostics11091538

Syahniar, R., Wahid, M. H., and Syam, A. F. (2019). Detecting the *Helicobacter pylori* 16S rRNA gene in dyspepsia patients using real-time PCR. *Acta Med Indones.* 51, 34–41.

Taborda, M. I., Aquea, G., Nilo, Y., Salvatierra, K., López, N., López, S., et al. (2018). Non-invasive diagnostic of *Helicobacter pylori* in stools by nested-qPCR. *Pol J Microbiol.* 67, 11–18. doi: 10.5604/01.3001.0011.5881

Wang, Y. K., Kuo, F. C., Liu, C. J., Wu, M. C., Shih, H. Y., Wang, S. S., et al. (2015). Diagnosis of *Helicobacter pylori* infection: current options and developments. *World J Gastroenterol.* 21, 11221-11235. doi: 10.3748/wjg.v21.i40. 11221

Yan, S., and Jun, Z. (2019). *Helicobacter pylori* recrudescence and its influencing factors. J Cell Mol Med. 23, 7919–7925. doi: 10.1111/jcmm.14682

Zamani, M., Masrour-Roudsari, J., and Zamani, V. (2017). Hematologic disorder: a manifestation of *helicobacter pylori* infection. *Caspian J Intern Med.* 8, 133–134. doi: 10.22088/cjim.8.2.133

Zhang, J., Cao, J., Zhu, M., Xu, M., and Shi, F. (2019). Loop-mediated isothermal amplification-lateral-flow dipstick (LAMP-LFD) to detect *Mycoplasma ovipneumoniae. World J. Microbiol. Biotechnol.* 35, 31. doi: 10.1007/s11274-019-2601-5