Rapid and Visual Detection of SARS-CoV-2 Using Multiplex Reverse Transcription Loop-Mediated Isothermal Amplification Linked With Gold Nanoparticle-Based Lateral Flow Biosensor

Xu Chen1,2,3*, Qingxue Zhou4†, Shijun Li3, Hao Yan5, Bingcheng Chang1,2, Yuexia Wang6 and Shilei Dong7*

1 The Second Clinical College, Guizhou University of Traditional Chinese Medicine, Guiyang, China, 2 Central Laboratory of the Second Affiliated Hospital, Guizhou University of Traditional Chinese Medicine, Guiyang, China, 3 Laboratory of Bacterial Infectious Disease of Experimental Centre, Guizhou Provincial Centre for Disease Control and Prevention, Guiyang, China, 4 Clinical Laboratory, Hangzhou Women’s Hospital, Hangzhou, China, 5 Department of Microbiology, Zhejiang Provincial Center for Disease Control and Prevention, Hangzhou, China, 6 TCM Research Institute, Zhejiang Chinese Medical University, Hangzhou, China, 7 Department of Clinical Laboratory, Zhejiang Hospital, Hangzhou, China

Background: Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is a novel coronavirus that has caused the outbreak of coronavirus disease 2019 (COVID-19) all over the world. In the absence of appropriate antiviral drugs or vaccines, developing a simple, rapid, and reliable assay for SARS-CoV-2 is necessary for the prevention and control of the COVID-19 transmission.

Methods: A novel molecular diagnosis technique, named multiplex reverse transcription loop-mediated isothermal amplification, that has been linked to a nanoparticle-based lateral flow biosensor (mRT-LAMP-LFB) was applied to detect SARS-CoV-2 based on the SARS-CoV-2 RdRp and N genes, and the mRT-LAMP products were analyzed using nanoparticle-based lateral flow biosensor. The mRT-LAMP-LFB amplification conditions, including the target RNA concentration, amplification temperature, and time were optimized. The sensitivity and specificity of the mRT-LAMP-LFB detection was 100%, and no cross-reactions to other respiratory pathogens were observed.

Results: The SARS-CoV-2 specific primers based on the RdRp and N genes were valid for the establishment of mRT-LAMP-LFB assay to detect the SARS-CoV-2 virus. The multiple-RT-LAMP amplification condition was optimized at 63°C for 30 min. The full process, including reaction preparation, viral RNA extraction, RT-LAMP, and product identification, could be achieved in 80 min. The limit of detection (LoD) of the mRT-LAMP-LFB technology was 20 copies per reaction. The specificity of mRT-LAMP-LFB detection was 100%, and no cross-reactions to other respiratory pathogens were observed.
INTRODUCTION

Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), a non-segmented positive-sense RNA genome virus, is a novel coronavirus that causes the outbreak of respiratory disease (COVID-19) all over the world (Bao et al., 2020; Zhang, 2020). In the 21st century, two important coronaviruses, severe acute respiratory syndrome coronavirus (SARS-CoV) and Middle East respiratory syndrome coronavirus (MERS-CoV), have severely threatened public health (in 2003 and 2012, respectively) (Chen, 2020; Wang et al., 2020). Since December 2019, the novel SARS-CoV-2 coronavirus has been found in many countries around the world and was declared as a disease of “public health emergency of international concern” by the World Health Organization (WHO) (Rothe et al., 2020). Most patients infected with SARS-CoV-2, present with acute onset of fever, cough, dyspnea, and radiological evidence of ground-glass lung opacities compatible with atypical pneumonia (Tu et al., 2020). Not only that, asymptomatic or mildly symptomatic cases have also been reported (Coronaviridae Study Group of the International Committee on Taxonomy of Viruses, 2020; Jiang et al., 2020). Owning to the current disease situation, the SARS-CoV-2 virus has been becoming the third coronavirus posing significant threats to public health worldwide. In the absence of appropriate antiviral drugs or vaccines, developing a reliable, simple, and rapid assay for SARS-CoV-2 is necessary for the prevention and control of the COVID-19 transmission.

The size of SARS-CoV-2 genome is ~30 kilobases and encodes ~9860 amino acids, which has been classified as a beta coronavirus (Ji, 2020; Younes et al., 2020). The genome of SARS-CoV-2 is arranged in the order of 5'-untranslated region (UTR), replicase complex (ORF1a/b), spike gene (S gene), E gene, M gene, N gene, 3' UTR, and several unidentified non-structural open reading frames (van Kasteren et al., 2020; Younes et al., 2020).

Conclusion: The mRT-LAMP-LFB technique developed in the current study is a simple, rapid, and reliable method with great specificity and sensitivity when it comes to identifying SARS-CoV-2 virus for prevention and control of the COVID-19 disease, especially in resource-constrained regions of the world.

Keywords: lateral flow biosensor, reverse transcription-loop-mediated isothermal amplification, limit of detection, COVID-19, SARS-CoV-2

MATERIALS AND METHODS

Materials Instruments

Viral RNA extraction kits (QIAamp Viral RNA minikits; Qiagen, Hilden, Germany) (Cat NO. 52906) were purchased from Qiagen (Beijing, China). Universal isothermal amplification kits, AMV Reverse Transcriptase, colorimetric indicator (malachite green, MG), and biotin-14-dCTP were obtained from Bei-Jing HaiTaiZhengYuan. Co., Ltd. (Beijing, China). The LFB
materials, including the backing card, sample pad, absorbent pad, conjugate pad, and nitrocellulose membrane (NC), were purchased from Jie-Yi Biotechnology, Co., Ltd. (Shanghai, China). Anti-FAM (rabbit anti-fluorescein antibody) and biotin-BSA (biotinylated bovine serum albumin) were purchased from Abcam. Co., Ltd. (Shanghai, China). Dye (Crimson red) streptavidin-coated polymer nanoparticles (129 nm, 10 mg ml⁻¹; 100 mM borate, pH 8.5, with 0.1% BSA, 0.05% Tween 20 and 10 mM EDTA) were purchased from Bangs Laboratories, Inc. (Indiana, USA).

**Design of RT-LAMP Primers**

Based on the reaction mechanism of LAMP, two sets of specific primers were designed according to the target genes RdRp and N (GenBank Accession No. NC_045512.2), respectively. The primers were designed with Primer Explorer V5 (http://primerexplorer.jp/e/; Eiken Chemical Co., Ltd., Tokyo, Japan) online primer design software and checked with the basic local alignment search tool (BLAST). The primer positions are shown in Figure 1, and the RdRp and N genes sequence alignment among seven human coronaviruses (SARS-CoV-2, SARS-CoV, MERS-CoV, HCoV-HKU-1, HCoV-NL63, HCoV-OC43, and HCoV-229E) are shown in Supplementary Figure 1. All of the primers were synthesized by TsingKe Biotech Co., Ltd. (Beijing, China) with HPLC purification grade.

**SARS-CoV-2 RNA Standard and Artificial SARS-CoV-2 Virus Preparation**

The SARS-CoV-2 RNA standard material was obtained from the Chinese Academy of Metrology (Code NO. GBW (E) 091089). The RNA transcripts contained ORF1ab gene segment (13201-15600), complete E gene, and N gene (GenBank NO. NC_045512), and the concentration of RNA was measured by absolute quantitative digital PCR.

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**FIGURE 1** | Sequence and location of the RdRp (A) and N (B) genes used to design SARS-CoV-2 mRT-LAMP primers. The nucleotide sequence of the sense strand of the RdRp and N is shown in the diagram. Right arrows and left arrows indicate sense and complementary sequences which were used in the current study, respectively.
TABLE 1 | The primers used in the present study.

| Primer name | Sequence and modifications | Length | Gene |
|-------------|---------------------------|--------|------|
| F3          | 5'-CACCTTATGGGTGGAGAT-3'  | 18 nt  | RdRp |
| B3          | 5'-AACATATAAGTAACCGCCA-3' | 19 nt  |      |
| FIP         | 5'-GCAAGAACAAGTAGGGCTAT-3' | 42 mer |
| BiP         | 5'-ACATACAGGTTGTTAGCCTGT-3' | 44 nt  |
| FIP*        | 5'-FAM-GCAAGAACAAGTAGGGCTAT-3' | 42 mer  |
| LF          | 5'-ATTCAAGTATGGCTGAA-3' | 19 nt  |
| LF*         | 5'-Biotin-ATTCAAGTATGGCTGAA-3' | 19 nt  |
| F3          | 5'-TGCTACATACAGGAAGCT-3' | 19 nt  | N    |
| B3          | 5'-TGCAAGATGTTGTAGGCTA-3' | 20 nt  |
| FIP         | 5'-TCTGGCCACGTCTATGATCGCTGCTGCTGAG-3' | 41 nt  |
| BiP         | 5'-AGACGGCATATGGTGTTGGC-3' | 40 nt  |
| FIP*        | 5'-Dg-TCTGGCCACGTCTATGATCGCTGCTGCTGAG-3' | 41 nt  |
| LF          | 5'-AAATACCATGTGGACGTGAGATC-3' | 23 nt  |
| LB          | 5'-AGGGAAGCTTGGATATACACCA-3' | 21 nt  |
| LF*         | 5'-Biotin-AAATACCATGTGGACGTGAGATC-3' | 23 nt  |

RdRp-FIP*, 5’-labeled with FAM when used in LAMP-LFB assay; RdRp-LF*, 5’-labeled with biotin when used in LAMP-LFB assay; N-FIP*, 5’-labeled with Dig when used in the LAMP-LFB assay; N*, 5’-labeled with biotin when used in the LAMP-LFB assay. FAM, 6-carboxy-fluorescein; Dig, digoxigenin; nt, nucleotide; mer, monomeric unit.

The pseudo-virus for the positive quality control agent was obtained from TsingKe Biotech Co., Ltd. (Beijing, China) (Cat NO. TSV2614), which was made with 293T cell cultures and included segments of the ORF1a/b gene (genome coordinates: 13237-13737, 15231-15729), M Gene (genome coordinates: 26523-27191), E Gene (genome coordinates: 26245-26472), and N Gene (genome coordinates: 28274-29533). The pseudo-virus of SARS-CoV and MERS-CoV were obtained from TsingKe Biotech Co., Ltd. (Beijing, China) (Cat NO. TSV2589; Cat NO. TSV2575).

RNA Template Preparation

In the current study, the viral RNA comes from both pseudo-virus (TsingKe Biotech Co., Ltd.) and clinical samples were obtained using Viral RNA Extraction Kits (Qiagen, Germany) in accordance with the manufacturer’s instructions. The RNA templates were stored at -80°C before use. The concentration was assayed using quantitative PCR with RNA standard. Then, 10-fold serial dilutions of the pseudo-viruses ranging from 1x10^4 copies/μl to 1 copies/μl were prepared.

Gold Nanoparticle-Based Lateral Flow Biosensor Preparation

The LFB platform was prepared according to a previous report (Cheng et al., 2019). Briefly, the LFB contained four components: an absorbent pad, NC membrane, sample pad, and conjugate pad (Jie-Yi Biotechnology, Co., Ltd.). The components were assembled orderly on a backing card. The capture reagents, including anti-FAM, anti-Dig, and biotin-BSA (Abcam, Co., Ltd.), were immobilized by physical adsorption on the reaction regions. Then, anti-FAM was immobilized at test line 1 (TL1) (RdRp), and anti-Dig was immobilized test line 2 (TL2) (N), while biotin-BSA was immobilized at the control line (CL); each line was separated by 5 mm. SA-PNPs (dye streptavidin-coated polymer nanoparticles) were gathered on the conjugate pad. The prepared biosensors were preserved in a plastic box with a desiccant gel at room temperature before use.

The Standard RT-LAMP Reaction

The single RT-LAMP reactions for RdRp or N were performed in 25 μl reaction systems as previously described. Briefly, 0.4 μM of each outer primer (F3 and B3), 0.8 μM of each loop primer (LF* and LB), 1.6 μM of each inner primer (FIP* and BIP), 0.4 mM of biotin-14-dCTP, 1 μl (8 U) of Bst DNA polymerase (New England Biolabs, USA), 1 μl (10 U) of AMV Reverse Transcriptase (New England Biolabs, USA), 12.5 μl of 2 × reaction buffer [40 mM Tris-HCl (pH 8.8), 40 mM of KCl, 16 mM of MgSO4, 20 mM of (NH4)2SO4, 2 M of betaine, and 0.2% Tween-20] (HuiDeXin Bio-technique, Tianjin, China), and 1 × 10^4 copies of the RNA template were added to a tube. The mixtures were incubated at 63°C for 1 h. Viral RNA from SARS-CoV (pseudo-virus), MERS-CoV (pseudo-virus), and double distilled water (DW) were used as negative controls (NCs). The mRT-LAMP reaction was performed in a one-step reaction in a 25 μl reaction system containing 12.5 μl of 2 × reaction buffer; 0.2 μM each outer primer, RdRp-F3, RdRp-B3, N-F3, and N-B3; 0.4 μM each loop primer, RdRp-LF*, RdRp-LB, N-LF* and N-LB; 0.8 μM each inner primer, RdRp-FIP*, RdRp-BIP, N-FIP* and N-BIP; 0.4 mM biotin-14-dCTP; 1 μl (8 U) of Bst DNA polymerase (New England Biolabs, USA); 1 μl (8 U) of AMV Reverse Transcriptase (New England Biolabs, USA); and 1 × 10^4 copies of RNA template. The reaction conditions were carried out as described above.

RT-LAMP Products Detection

The monitoring techniques, including 2% agarose gel electrophoresis, visual detection reagents MG (VDR, Haitai-Zhengyuan biotech, Co. Ltd. Beijing, China), and lateral flow biosensor (LFB) methods, were applied for the determination and verification of the RdRp-RT-LAMP, N-RT-LAMP, and mRT-LAMP products. For the products amplified effectively, the agarose gel presented ladder-like bands, and the color changed from colorless to light green in the MG assay. However, there have no bands in gel electrophoresis, and the
color remains colorless in negative and blank controls. The strategy of visualization of RT-LAMP products with LFB was as previously described (Gong et al., 2019).

**Temperature Optimization of the RT-LAMP Assays**

To confirm the optimal amplification temperature for RdRp-RT-LAMP and N-RT-LAMP, the pseudo-virus of SARS-CoV-2- RdRp-N was used as a positive control at a concentration of 1×10^4 copies per reaction, and the RT-LAMP amplifications were monitored by a real-time turbidity technique. Reaction temperatures ranging from 60 to 67°C with 1°C intervals were tested. The curves of DNA concentrations of each amplified product were exhibited in the graph. Turbidity > 0.1 was considered as positive. Three replicates were tested for each temperature.

**Optimization of the Amplification Time for the mRT-LAMP-LFB Assay**

To optimize the reaction time of mRT-LAMP-LFB, four amplification times (20, 30, 40, and 50 min) were evaluated. The mRT-LAMP-LFB reactions were carried out as described above, and the results were tested by LFB. Each reaction time was tested at least three times.

**Analytical Sensitivity of mRT-LAMP-LFB Assays**

The sensitivity of each RT-LAMP-LFB reaction (RdRp-RT-LAMP-LFB, N-RT-LAMP-LFB, and mRT-LAMP-LFB) was determined using pseudo-virus of SARS-CoV-2 with ten-fold serial dilutions range from 1×10^4 copies to 1 copy. The RT-LAMP reactions were carried out as described above, and the results were tested using visual detection reagents (MG) and LFB. The limit of detection (LoD) of single and multiplex reactions was verified as the last dilution of each positive test. The LoD of RT-PCR technology using Applied Biosystems™ 7500 Real-Time PCR System (Life Technologies, Singapore) with Novel Coronavirus Nucleic Acid Diagnostic Real-Time RT-PCR Kit (Sansure biotech Inc, China) was used as the reference standard, which was recommended by the Chinese Center for Disease Control and Prevention. The RT-PCR detection was performed with Applied Biosystems™ 7500 Real-Time PCR System (Life Technologies, Singapore). A threshold cycle (Ct) value < 38 was determined to indicate a positive result. The mRT-LAMP-LFB and RT-PCR assays were performed simultaneously in a biosafety level 2 laboratory, as detailed in the WHO Laboratory biosafety manual, third edition. The mRT-LAMP-LFB detection was performed as described above.

**RESULTS**

COVID-19 is a newly emerging, life-threatening respiratory disease caused by a novel coronavirus SARS-CoV-2, and it has had a significant impact on public health and the economy worldwide (Bao et al., 2020; She et al., 2020). The purpose of the current study is to develop a reliable, rapid, sensitive, and easy-to-use assay for SARS-CoV-2.

**Verification and Analysis of RT-LAMP Products**

To confirm the amplification with the two sets of LAMP primers, the RdRp-, N-, or mRT-LAMP mixtures were incubated at a constant temperature of 65°C for 1 h. Then, the RdRp-, N-, and mRT-LAMP products were analyzed with 2% agarose gel electrophoresis, colorimetric indicator (MG), and lateral flow biosensor (LFB), respectively. The ladder-like bands of agarose gel were observed in the positive amplification, but not in the negative controls (Figures 2A, D, G). The color of the positive results in the RdRp-, N-, and mRT-LAMP reactions changed from colorlessness to bright green, while the negative reactions remained colorless (Figures 2B, E, H). LFB was used for further confirmation of RdRp-, N-, and mRT-LAMP. For RdRp-RT-LAMP detection, two crimson red bands (CL and TL1) appeared, indicating positive results. CL and TL2 were visible for N-RT-LAMP, indicating successful amplification, while the negative controls only appeared as a crimson red line (CL) in the biosensor (Figures 2C, F, I). Therefore, the results suggested that the two sets of RT-LAMP primers for RdRp and N detection were valid for the development of the mRT-LAMP assay.

**Application of the mRT-LAMP-LFB Method to Analyze the Clinical Samples and Artificial Sputum Samples**

To verify the applicability of the mRT-LAMP-LFB assay for detecting SARS-CoV-2, one hundred and ten clinical nasopharyngeal swab specimens were collected from suspected SARS-CoV-2-infected patients, and sixty artificial sputum samples (randomly added 100 copies of SARS-CoV-2 pseudo-viruses in each 200 µl artificial sputum sample) were used in the current study. The artificial sputum samples were pretreated with N-acetyl-L-cysteine-2% NaOH. The initial process of all specimens was handled in a validated biological safety cabinet, and performed by staff trained with appropriate personal protective equipment. The clinical samples and artificial sputum samples were detected for SARS-CoV-2 using RT-PCR and mRT-LAMP-LFB methods. The mRT-LAMP detection was as described above. The Novel Coronavirus Nucleic Acid Diagnostic Real-Time RT-PCR Kit (Sansure biotech Inc, China) was used as the reference standard, which was recommended by the Chinese Center for Disease Control and Prevention. The RT-PCR detection was performed with Applied Biosystems™ 7500 Real-Time PCR System (Life Technologies, Singapore). A threshold cycle (Ct) value < 38 was determined to indicate a positive result. The mRT-LAMP-LFB and RT-PCR assays were performed simultaneously in a biosafety level 2 laboratory, as detailed in the WHO Laboratory biosafety manual, third edition. The mRT-LAMP-LFB detection was performed as described above.
amplification was tested at different temperatures (60 to 67°C with 1°C intervals) with genomic templates (1×10⁴ copies) from the pseudo-virus of SARS-CoV-2. The RT-LAMP amplification protocol was as described above, the RdRp- and N-LAMP amplifications were monitored by means of real-time turbidity technique, and the kinetics graphs were recorded from all temperatures. The results showed that the faster amplifications of RdRp-RT-LAMP were obtained for detection temperature range from 63 to 64°C, and 62 to 63°C for the N-RT-LAMP reactions (Figure 3). Hence, the amplification temperature of

### TABLE 2 | Pathogens used in the current study.

| No. | Pathogen species | Pathogen name | Source of pathogens | No. of strains | RT-LAMP-LFB result⁵ \\  |  |  |
|-----|------------------|---------------|---------------------|---------------|------------------|
|     | Coronavirus      |               |                     |               | RdRp        | N |
| 1   | SARS-CoV-2 (pseudo-virus) | 2019-nCoV-ab II EMN | TsingKe Biotec Co., Ltd. | 1 | P | P |
| 2   | SARS-CoV-2 (nucleic acid samples) | ZJCDC-2019-nCoV-52; -85; -86; -90; -120; -134; -152; -189; -190; ZJ-2019-nCoV-304; -305 | TsingKe Biotec Co., Ltd. (Beijing, China); ZJCDC and 1st ZJUSM | 12 | P | P |
| 3   | SARS-CoV (pseudo-virus) | SARS-ORF1a-N | TsingKe Biotec Co., Ltd. (Beijing, China) | 1 | N | N |
| 4   | MERS-CoV (pseudo-virus) | MERS-abEN | TsingKe Biotec Co., Ltd. (Beijing, China) | 1 | N | N |
| 5   | Human coronavirus HKU1 | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 6   | Human coronavirus HCoV-NL63 | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 7   | Human coronavirus OC43 | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 8   | Human coronavirus 229E | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
|     | Other pathogens  |               |                     |               | RdRp        | N |
| 9   | H1N1 | ZJH-H1N1-57 | Zhejiang Hospital | 1 | N | N |
| 10  | H3N2 (nucleic acid sample) | GZCDC-H3N2-14 | GZCDC | 1 | N | N |
| 11  | H5N1 (nucleic acid sample) | GZCDC-11-H5N1 | GZCDC | 1 | N | N |
| 12  | H7N9 (nucleic acid sample) | GZCDC-5-H7N9 | GZCDC | 1 | N | N |
| 13  | Influenza B | ZJH-Influenza-B-115 | Zhejiang Hospital | 1 | N | N |
| 14  | Respiratory syncytial virus type A | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 15  | Respiratory syncytial virus type B | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 16  | Human rhinovirus | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 17  | Adenoviruses | Quality control sample | DAAN Gene Co., Ltd. (Guangzhou, China) | 1 | N | N |
| 18  | Mycoplasma pneumoniae | ZJH-MP-594 | Zhejiang Hospital | 1 | N | N |
| 19  | Mycobacterium tuberculosis | GZCDC-MTB-564 | GZCDC | 1 | N | N |
| 20  | Pseudomonas aeruginosa | ATCC 27853 | ATCC | 1 | N | N |
| 21  | Klebsiella pneumonia | ZJH-KP-104 | Zhejiang Hospital | 1 | N | N |
| 22  | Streptococcus pneumoniae | ZJH-SP-016 | Zhejiang Hospital | 1 | N | N |
| 23  | Mycoplasma pneumoniae | GZUTOM-MP-102 | 2nd GZUTOM | 1 | N | N |
| 24  | Haemophilus influenzae | ATCC49247 | ATCC | 1 | N | N |
| 25  | Streptococcus pyogenes | ZJH-SP-1087 | Zhejiang Hospital | 1 | N | N |
| 26  | Acinetobacter baumannii | ZJH-AB-984 | Zhejiang Hospital | 1 | N | N |
| 27  | Staphylococcus aureus | ZJH-SA-065 | Zhejiang Hospital | 1 | N | N |
| 28  | Cryptococcus neoformans | ATCC14053 | ATCC | 1 | N | N |
| 29  | Candida glabrata | ZJH-CG-057 | Zhejiang Hospital | 1 | N | N |
| 30  | Hemophilus parainfluenza | GZCDC-HP-045 | GZCDC | 1 | N | N |
| 31  | Shigella boydii | GZCDC-SB-107 | GZCDC | 1 | N | N |
| 32  | Enteropathogenic Escherichia coli | GZCDC-EPEC-045 | GZCDC | 1 | N | N |
| 33  | Bordetella pertussis | GZCDC2-BP-052 | GZCDC | 1 | N | N |
| 34  | Bordetella parapertussis | GZCDC2-BP-0094 | GZCDC | 1 | N | N |
| 35  | Bacillus cereus | GZUTOM-BC-037 | GZUTOM | 1 | N | N |
| 36  | Listeria monocytogenes | GZUTOM-LM-025 | GZUTOM | 1 | N | N |
| 37  | Neisseria flexneri | GZUTOM-SF-018 | GZUTOM | 1 | N | N |
| 38  | Leptospira interrogans | GZCDC-LJ-005 | GZCDC | 1 | N | N |

| a | ZJCDC, Zhejiang Provincial Center for Disease Control and Prevention; 1st ZJUSM, The First Affiliated Hospital, Zhejiang University School of Medicine; ZJCCL, Zhejiang Center for Clinical Laboratories; 2nd GZUTCM, The Second Affiliated Hospital, Guizhou University of Traditional Chinese Medicine; GZCDC, Guizhou Provincial Center for Disease Control and Prevention; ATCC, American Type Culture Collection.  
| b | P, Positive; N, Negative. |
63°C was considered as optimal temperature for the rest of multiple-RT-LAMP reactions in the current study.

**Optimization of Amplification Time for mRT-LAMP-LFB Assay**
To obtain an optimal reaction time for mRT-LAMP, four amplification times (20, 30, 40, and 50 min) were tested at the 63°C amplification temperature. The results showed that the LoD of the genomic RNA templates (20 copies) was detected when the mRT-LAMP amplification lasted 30 min (Figure 4). Hence, a reaction time of 30 min was considered the optimal amplification time for mRT-LAMP detection. In summary, the whole detection procedure, including reaction preparation (approximately 10 min), target genomic RNA preparation (30 min), mRT-LAMP (30 min), and analysis of results (approximately 2 min), could be completed within 80 min.

**Sensitivity of RdRp-, N-, and mRT-LAMP Detection**
The sensitivity of RdRp-, N-, and mRT-LAMP detection was evaluated with serially diluted pseudo-virus RNA range from 1×10⁴ copies to 1 copy. The RT-LAMP amplification products were analyzed by visual inspection with MG reagents and lateral flow biosensors. The CL and TL1 lines appeared on the biosensor, showing positive results for the RdRp-RT-LAMP assay, and two crimson lines (CL and TL2) were observed on the biosensor, indicating positive results for N-RT-LAMP detection. The CL, TL1, and TL2 bands simultaneously became crimson on the biosensor, reporting positive results for the RdRp and N genes. For the negative controls, only the CL line appeared on the biosensors. The results showed that the LoD of mRT-LAMP was 20 copies per reaction, which was the same as the LoD of the RdRp- and N-RT-LAMP assay (Figures 5A, B, D, E, G, H). Meanwhile, the sensitivity of RT-PCR technique was also tested in the current study, the results indicated that the LoD of RT-PCR was 100 copies per reaction (Figures 5C, F, I).

**Specificity of the mRT-LAMP Assay**
The specificity of mRT-LAMP detection was confirmed with pseudo-viruses of SARS-CoV-2, 12 clinical SARS-CoV-2-positive samples, and 36 other pathogens (Table 2). The process of mRT-LAMP amplification, as described above. The genomic RNA extracted from SARS-CoV-2 presented positive results. Other pathogens and the blank control showed negative results (Table 2). Hence, the results confirmed that the mRT-
LAMP-LFB method could accurately identify SARS-CoV-2 from other pathogens.

**Feasibility of the mRT-LAMP-LFB Method Using Clinical Samples**
To further demonstrate the feasibility of mRT-LAMP-LFB as a valuable method for the detection of SARS-CoV-2, 110 clinical nasopharyngeal swab specimens and 60 artificial sputum samples (randomly added 100 copies of SARS-CoV-2 pseudoviruses in each 200 μL artificial sputum sample) were simultaneously tested by mRT-LAMP-LFB and RT-PCR. Among them, 12 clinical samples and 35 artificial sputum samples had been confirmed as SARS-CoV-2 through RT-PCR and mRT-LAMP-LFB, respectively (Table 3). The Cq values of RT-PCR and mRT-LAMP-LFB detection results were shown in Supplementary Table 1. These results suggested that the mRT-LAMP-LFB assay established in the current study could be used as an advanced tool to detect SARS-CoV-2.

**DISCUSSION**
SARS-CoV-2 is the seventh coronavirus that causes human infections. Like SARS-CoV and MERS-CoV, this virus has the ability to cause lethal pneumonia (Chiappelli, 2020). Moreover, it has a stronger human-to-human transmission capacity than the above two coronaviruses (Ki, 2020; Wilson and Chen, 2020). Until now, up to 140 million COVID-19 cases have been confirmed, including more than 3 million deaths (www.who.int/emergencies/diseases/novel-coronavirus-2019).
The main findings of the current study are that we established a simple, sensitive, reliable, and rapid assay with great specificity and low equipment cost for SARS-CoV-2 by mRT-LAMP-LFB. To avoid false-positive or -negative results, we chose the two target genes, RdRp and N, to detect viral RNA in clinical samples (Chen et al., 2020; Huang et al., 2020; Pang et al., 2020). To reduce the amplification time, we designed the loop primers. Briefly, six primers targeting eight regions generated a self-priming dumbbell-shaped template upon isothermal incubation with strand-displacing polymerase, resulting in the

**FIGURE 4** Optimization of the amplification time for mRT-LAMP-LFB detection. Different amplification times (A, 20 min, B, 30 min, C, 40 min, D, 50 min) were tested at 63°C. Biosensors 1-7 represent SARS-CoV-2 genomic RNA levels of 1×10^4 copies, 1×10^3 copies, 1×10^2 copies, 20 copies, 10 copies, and 1 copy per reaction and blank control (DW), respectively. The best sensitivity was observed when the amplification lasted for 30 min (B).

**FIGURE 5** Sensitivity analysis of RdRp-, N-, and mRT-LAMP detection with serial dilutions of RNA extracted from pseudo-virus SARS-CoV-2. The LoD of RT-LAMP for detecting SARS-CoV-2 was analyzed with visual detection reagents (MG) and lateral flow biosensors. (A, B) Sensitivity analysis of RdRp-RT-LAMP reaction. Tubes A1-A7 (Biosensors B1-B7) represent the genomic RNA amounts of 1×10^4 copies, 1×10^3 copies, 1×10^2 copies, 20 copies, 10 copies, and 1 copy per reaction and blank control (DW), respectively. The LoD of RdRp-RT-LAMP detection was 100 copies of RNA template per reaction. (D, E) Sensitivity analysis of N-RT-LAMP reaction. Tubes D1-D7 (Biosensors E1-E7) represent the genomic RNA amounts of 1×10^4 copies, 1×10^3 copies, 1×10^2 copies, 20 copies, 10 copies, and 1 copy per reaction and blank control (DW), respectively. The LoD of N-RT-LAMP detection was 20 copies of RNA template per reaction. (G, H) Tubes G1-G7 (Biosensors H1-H7) represent the genomic RNA amounts of 1×10^4 copies, 1×10^3 copies, 1×10^2 copies, 20 copies, 10 copies, and 1 copy per reaction and blank control (DW), respectively. The LoD of the mRT-LAMP assay for RdRp and N detection was 20 copies of RNA template per reaction. (I) Sensitivity of mRT-PCR detection (1×10^4 copies to 1 copy). The LoD of mRT-PCR detection was 100 copies of RNA template per reaction.
In conclusion, a simple, rapid, and reliable mRT-LAMP-LFB technique based on the RdRp and N genes was successfully developed for assaying SARS-CoV-2 in the current study. This method could rapidly, reliably, specifically, and sensitively detect SARS-CoV-2. The amplification products were analyzed with LFB, which was objective, rapid, and easily interpretable. Hence, the mRT-LAMP-LFB assay could be considered as a useful method for the reliable and rapid detection of SARS-CoV-2 in clinical samples, especially in resource-constrained regions of the world.

### DATA AVAILABILITY STATEMENT

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

### ETHICS STATEMENT

The study was approved by the Human Ethics Committee of the Second Affiliated Hospital of Guizhou University of Traditional Chinese Medicine (Approval No. TYH2020011) and the Human Ethics Committee of the Zhejiang Hospital (Approval No. 202010717954. X). The shortcoming of this detection is that the RT-LAMP amplification must be taken out from the reaction tube for LFB detection. There is a risk of contamination with the post-reaction processing of LAMP products. The strict control of the laboratory environment is critical for the reduction of the production of aerosols in experimental processes. Spraying timely 10~15% sodium hypochlorite solution and 70% ethanol after completion of detection is an effective way to overcome nucleic acid contamination in the laboratory. In the current study, the mRT-LAMP-LFB detection results were consistent with the RT-PCR methods in the evaluation of clinical samples. It is indicated that false-positive rates have been effective controlled in our laboratory.

The main limitation of this study is that with the widely spread of SARS-CoV-2 virus, the accuracy of the mRT-LAMP-LFB technology will be affected by the mutations occurring in the primers sequence region of the target genes. So, it is necessary to monitor the mutant sites of the virus genome by whole-genome sequencing. Besides, owning to laboratory biosafety, SARS-CoV and MERS-CoV viruses could not be tested for the specificity of the mRT-LAMP-LFB assay, we used pseudo-virus of SARS-CoV and MERS-CoV as alternatives.

In conclusion, a simple, rapid, and reliable mRT-LAMP-LFB technique based on the RdRp and N genes was successfully developed for assaying SARS-CoV-2 in the current study. This method could rapidly, reliably, specifically, and sensitively detect SARS-CoV-2. The amplification products were analyzed with LFB, which was objective, rapid, and easily interpretable. Hence, the mRT-LAMP-LFB assay could be considered as a useful method for the reliable and rapid detection of SARS-CoV-2 in clinical samples, especially in resource-constrained regions of the world.

### TABLE 3 | Comparison of RT-PCR and mRT-LAMP-LFB methods to identify SARS-CoV-2 in clinical samples and artificial sputum samples.

| Detection method  | Clinical samples (n = 110) | Artificial sputum samples (n = 60) |
|-------------------|----------------------------|-----------------------------------|
|                   | Positive | Negative | Time consumption | Positive | Negative | Time consumption |
| RT-PCR            | 12 (Ct<38)| 98       | ~150 min        | 35 (Ct<38)| 25       | ~150 min        |
| mRT-LAMP-LFB      | 12       | 98       | Within 80 min   | 25       | 25       | Within 80 min   |

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In conclusion, a simple, rapid, and reliable mRT-LAMP-LFB technique based on the RdRp and N genes was successfully developed for assaying SARS-CoV-2 in the current study. This method could rapidly, reliably, specifically, and sensitively detect SARS-CoV-2. The amplification products were analyzed with LFB, which was objective, rapid, and easily interpretable. Hence, the mRT-LAMP-LFB assay could be considered as a useful method for the reliable and rapid detection of SARS-CoV-2 in clinical samples, especially in resource-constrained regions of the world.

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Lin Shen Di (7K) Hao, and complied with the Declaration of Helsinki. All data/isolates were isolated anonymously.

AUTHOR CONTRIBUTIONS

XC, QZ, and SD conceived and designed the study. XC and SD participated in primers design. XC, QZ, BC, YW, and HY contributed to all the laboratory works. BC and HY contributed to the data collection. XC, SL, and QZ performed the statistical analysis. XC wrote the initial draft of the manuscript, and SD revised the manuscript. All authors contributed to the article and approved the submitted version.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fcimb.2021.581239/full#supplementary-material

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