Published online 2020 July 25.

Research Article

Diagnostic Values of Multiplex Loop-Mediated Isothermal Amplification and Multiplex Polymerase Chain Reaction for Detection of Methicillin-Resistant *Staphylococcus aureus*

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Received 2019 July 29; Revised 2020 June 17; Accepted 2020 June 17.

Abstract

Background: Methicillin-resistant *Staphylococcus aureus* (MRSA) has emerged as a significant pathogen in community and hospital environments and is associated with high mortality and morbidity. Both polymerase chain reaction (PCR) and loop-mediated isothermal amplification (LAMP) methods are sensitive and acceptable molecular methods for the diagnosis of infectious diseases. **Objectives:** This study aimed to develop detection assays for Staphylococcal *mecA* and *spa* using multiplex PCR and LAMP.

Methods: Both methods were standardized, and detection limits were determined using serial dilutions of *S. aureus* DNA samples. Fifty-three clinical isolates of *S. aureus* were confirmed to the species level using biochemical tests and multiplex PCR and multiplex LAMP for the *spa* gene, while disk diffusion, minimum inhibitory concentration, and detection of *mecA* genes were used for the assessment of methicillin resistance.

Results: The PCR could detect the *mecA* and *spa* genes at 1 fg/mL and 10 fg/mL of bacterial DNA, which were equal to 35 and 350 gene copy numbers, respectively. Similarly, multiplex LAMP detected the *spa* and *mecA* genes at 0.1 fg/mL and 1 fg/mL of bacterial DNA, which were equal to 3.5 and 35 genome copy numbers, respectively. According to MIC and disk diffusion methods, four (7.54%) cases were oxacillin-sensitive methicillin-resistant *S. aureus*, 16 isolates were methicillin-sensitive, and 37 isolates were methicillin-resistant. According to multiplex PCR, 47.75% of the isolates were *mecA*-positive while in multiplex LAMP, 41 (35.77%) isolates were *mecA*-positive.

Conclusions: The sensitivity and specificity of the multiplex LAMP were higher than those of multiplex PCR and biochemical methods. Thus, we can apply the LAMP for the routine detection of MRSA.

Keywords: Staphylococcus aureus, Molecular Diagnosis, MRSA, Multiplex LAMP, Multiplex PCR

1. Background

Staphylococcus aureus is a commensal bacterium that usually colonizes the skin and mucous membranes and causes a wide variety of diseases, including soft-tissue and bloodstream infections and life-threatening sepsis (1, 2). Peptidoglycan biosynthesis in the cell wall of Grampositive bacteria starts with membrane-bound enzymes called penicillin-binding proteins (e.g., PBP2). Due to the acquisition of *mecA* in certain strains, a new protein called PBP2a is encoded. This protein cannot attach to betalactam antibiotics, although it retains the enzymatic activity; thus, these strains are called methicillin-resistant *S. aureus* (MRSA) (3).

In recent years, the frequency of MRSA has increased to 60% of the clinical isolates of *S. aureus* (4, 5). Some

clinical *mecA* and PBP2 positive strains of *S. aureus* exhibit phenotypic susceptibility to oxacillin. It has been suggested that such isolates can be classified as a new group of MRSA called oxacillin-susceptible methicillin-resistant *S. aureus* (OS-MRSA). They may be incorrectly classified as methicillin-sensitive *S. aureus* (MSSA) just because they show phenotypic susceptibility to oxacillin (6, 7). It is estimated that the prevalence of MRSA and the mortality rate of *S. aureus* bacteremia in many countries have raised to more than 49% and 50%, respectively (8, 9). Recently, the epidemiology of MRSA has changed, and several nosocomial strains have produced community-acquired infections, named community-onset hospital-acquired infections (CO-HA MRSA), or vice versa (HO-CA MRSA) (10, 11).

The *mecA* gene is carried by the cassette chromosome *mecA* (SCCmec). Eleven *mecA* cassettes (SCCmec) are rec-

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ognized as *S. aureus* (12, 13). The majority of the prevalent MRSA strains in the community are *SCCmec* type IV, while types I, II, and III are associated with hospital-acquired MRSA infections (5, 14). The routine detection methods of MRSA, like culture and PCR, have several strengths and weaknesses. The disadvantages of these techniques include the high cost of the instruments, and more importantly, the inability to apply as a point of care test (15, 16). Thus, the development of a low-cost and fast method with high sensitivity and specificity is of utmost importance (17, 18).

As an alternative method for the diagnosis of MRSA, loop-mediated isothermal amplification (LAMP) was developed by Notomi et al. (19) in 2000. This method uses the *Bst* DNA polymerase that has the ability of strand displacement and yields large fragments of DNA by the autocycling reaction. In addition, the LAMP reaction is performed at 60° C - 65° C and is carried out for 60 to 90 min (20, 21). This method involves six primers especially designed to recognize a total of six distinct sequences on the target DNA. The LAMP method has been used to identify many bacteria, such as *Helicobacter pylori* and *Arcobacter*, and many parasites, fungi, and viruses (1, 6, 21, 22).

2. Objectives

Concerning the high prevalence of the infections caused by MRSA strains, especially in immunecompromised patients, and the importance of this bacterium in nosocomial infections, we attempted to use LAMP for the diagnosis of MRSA in clinical settings.

3. Methods

3.1. Bacterial DNA

Serial dilutions of DNA were prepared from a confirmed clinical isolate of *S. aureus*. Bacterial DNA was extracted from a 24-h cultured organism using the CinnaPure-DNA kit following the manufacturer's protocol.

3.2. Clinical Sample

During 2013 to 2014, 53 suspected clinical samples of MRSA and MSSA isolated at the Microbiology Laboratory of Imam Reza Hospital, Kermanshah, Iran, were re-identified to the species level using standard biochemical tests like Gram staining, catalase, DNase, coagulase, oxidase, and sensitivity to bacitracin (23).

3.3. Antimicrobial Susceptibility Testing

The disk diffusion and mico-dilution broth methods were carried out according to the standard methods using cefoxitin and oxacillin disks (MAS diagnosis) and oxacillin powder (Sigma), respectively. The in vitro antimicrobial susceptibility testing for MRSA was performed using the Kirby-Bauer disk diffusion method (Clinical Laboratory and Standard Institute, CLSI). The MIC of oxacillin was determined using serial dilutions from 0.5 to 1,024 μ g/mL of the antibiotic (CLSI, 2016) (24).

3.4. PCR and LAMP Primer Design

The sets of forward and reverse PCR primers for *16sRNA*, *mecA*, and *spa* genes were designed manually and confirmed by the online Primer3 software (Table 1). The LAMP primers including outer forward primer (F3), outer backward primer (B3), forward inner primer (FIP), and backward inner primer (BIP) targeting the *spa* and *mecA* genes were designed by online LAMP primer design software Primer Explorer V4 (Table 2).

Table 1. Polymerase Chain Reaction Primer Sequences		
Target Gene	Primer (5' $ ightarrow$ 3')	Size of Amplicon
spa	F: CTAATAACGCTGCACCTAAG	309
	R: GCAAAGAAGATGGCAACAAAC	
mecA	F: ACCACTTCATATCTTGTAACG	160
	R: AGATTACAACTTCACCAGGTTC	
16srRNA	F: GCAAGCGTTATCCGGATTT	- 597
	R: CTTAATGATGGCAACTAAGC	

Table 2. Loop-mediated Isothermal Amplification Primer Sequences

Target Gene	Primer $(5' \rightarrow 3')$
spaF3	TTATAGTTCGCGACGACG
spaB3	TGCAGATAACAAATTAGCTGAT
spaFIP	CCAGAAACTGGTGAAGAAAATCCATGAATTCCTAAGGCTAATGATAATCCACC
spaBIP	AATGCTTGAGCTTTGTTAGCATCT-GAATTC-AACATGATCAAACCTGGTCA
spaLF	TCATCGGTACAACTGTA
spaBF	GCATGGTTTGCTGGTTGC
mecAF3	AGTTCTTTAGCGATTGCTTTA
mecAB3	CATTGATCGCAACGTTCAA
mecAFIP	CGAAACAATGTGGAATTGGCCAA-GAATTC-ATTCTTTGGAACGATGCCTAT
mecABIP	GCTTTGGTCTTTCTGCATTCCTG-GAATTC-AGAAGATGGTATGTGGAAGT
mecALF	ACAGGAACAGCATATGAG
mecABF	ATGACGCTATGATCCCAA

3.5. Molecular Methods

3.5.1. Multiplex PCR

First, the uniplex PCR was set up for 16sRNA, spa, and mecA separately. The PCR mixtures contained 1.5 μ L 10x PCR

buffer, 1.5 mmol MgCl₂, 0.2 mmol dNTP, 1 μ mol of each of the designed primers, 1 U Taq DNA polymerase, 2.0 μ L template DNA, and 8.55 µL distilled H₂O to a final volume of 15.0 μ L. The optimum thermal cycling program for all the gene targets was set as follows: initial denaturation at 95°C for 5 min, followed by 34 cycles of denaturation at 95°C for 30 s, annealing at 53°C for 30 s, extension at 72°C for one minute, and a final extension at 72°C for 5 min. As we aimed to identify the spa and mecA genes simultaneously, the designed program was used to amplify the genes at the same time in a multiplex reaction. Therefore, the PCR mixture was prepared as follows: 2 μ L 10 × PCR buffer, 0.6 μ L MgCl₂, 0.5 μ L dNTP, 2 μ L each primer, 0.3 μ L Taq DNA polymerase, 2.0 μ L template DNA, and 5.6 μ L H₂O to a final volume of 15.0 μ L. The program was set as mentioned before, except that the annealing temperature was set at 54°C for both genes. Finally, the PCR products were detected using electrophoresis on a 1.5% agarose gel using ethidium bromide staining.

3.5.2. Multiplex LAMP

The uniplex LAMP was set up for the spa and mecA genes before designing multiplex LAMP. The LAMP mixture was set as follows: 2 μ L 10 \times buffer, 2.5 μ L MgSO₄, 5 μ L dNTP, 4 μ L Betine, 0.4 μ L of each of FIP and BIP primers, 0.5 μ L of each of LF and LB primers, 2 μ L template DNA, 1.1 μ L Bst polymerase, and 4.3 μ L H₂O to a final volume of 25 μ L. This program was set as follows: initial denaturation at 92°C for 5 min, 60.5 $^{\circ}\text{C}$ for 40 min, and a final extension at 92 $^{\circ}\text{C}$ for 3 min. The multiplex LAMP reaction for the spa and mecA genes was carried out on a scale of 30 μ L mixture containing 6 μ L 10× LAMP buffer (1.5×), 3.2 mmol (8 μ L) dNTP, 0.8 μ mol betaine, and 8 mmol MgSO₄. For the *mecA* gene, 1.2 μ mol/L of primers FIP and BIP, 0.16 μ mol/L of primers F3 and B3, and 0.4 μ mol/L of primers LF and LB were applied. For the spa gene, 1.6 μ mol/L of primers FIP and BIP, 0.2 μ mol/L of primer F3, 0.16 μ mol/L of primer B3, and 0.8 μ mol/L of primers LF and LB were mixed. Besides, 2 μ L of template DNA and 4.2 μ L of sterile deionized water was used for all the mixtures and were incubated at 92°C for 5 min. After this step, 1.7 μ L of Bst DNA polymerase was added and incubated at 60.5°C for 80 min, heated to 92°C for 3 min, and finally cooled at 4°C for 5 min to terminate the reaction. For further confirmation, the products were also detected using electrophoresis on a 1.5% agarose gel using ethidium bromide staining.

3.6. Analytical Sensitivity of PCR and LAMP

The analytical sensitivity of LAMP and PCR was determined Using nanodrop instrument, the amount of DNA in the first concentration was determined as 83.3 mg/mg. Then, different concentrations of pure DNA were prepared by serial dilutions.

3.7. Analytical Specificity of LAMP and PCR

To determine the analytical specificity of designed LAMP and PCR for the *mecA* and *spa* genes, the total genomic DNA of methicillin-sensitive *S. epidermidis*, *Klebsiella*, *Acinetobacter*, *Escherichia coli*, *Citrobacter freundii*, *Enterococcus faecalis*, *Micrococcus*, *S. agalactiae*, *S. saprophyticus*, and *Enterobacter* was tested by the designed methods.

3.8. Clinical Sensitivity and Specificity

Fifty-three isolates of suspected *S. aureus* were determined by biochemical methods and confirmed by the amplification of the *16S rRNA* gene. Resistance to methicillin was surveyed by disk diffusion using cefoxitin and oxacillin disks; the MIC of oxacillin was determined, as well. To determine the clinical sensitivity and specificity of PCR and LAMP, these tests were compared with Gold standards, including culture, antibiogram, and MIC.

3.9. Statistical Analysts

The data from LAMP and PCR methods were analyzed using SPSS version 21. The agreement between the two methods was analyzed by the kappa coefficient test.

4. Results

4.1. Analytical sensitivity of LAMP and PCR

The analytical sensitivity of multiplex PCR and multiplex LAMP was determined using serial dilutions of *S. aureus* DNA samples as the target DNA. The PCR could detect the *mecA* and *spa* genes at 1 fg/mL and 0.01 pg/mL of bacterial DNA, which were equal to 35 and 350 copy numbers of bacterial DNA, respectively. Similarly, multiplex LAMP detected the *spa* and *mecA* genes at 0.1 fg/mL and 1 fg/mL of bacterial DNA, which were equal to 3.5 and 35 copy numbers of bacterial DNA, respectively (Figures 1 and 2).

4.2. Analytical Specificity of LAMP and PCR

Multiplex LAMP and multiplex PCR were carried out on methicillin-sensitive *S. epidermidis*, *Klebsiella* sp., *Acinetobacter* sp., *E. coli*, *Citrobacter* sp., *E. faecalis*, *Micrococcus* sp., *S. agalactiae*, *S. saprophyticus*, and *Enterobacter* sp. Amplification was not observed in both tests, which showed a 100% analytical specificity of the tests.

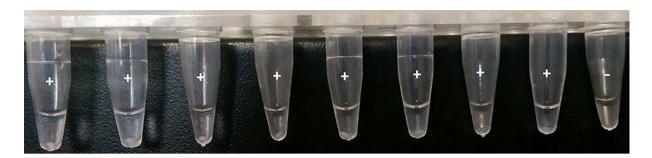


Figure 1. Agarose gel electrophoresis of multiplex LAMP products of spa and mecA genes. Lane 1: mecA positive control, lane 2: spa positive control, A, 1 ng/µL of bacterial DNA; B, 0.1 ng/µL; C, 0.01 ng/µL; D, 1 pg/µL; E, 0.1 pg/µL; F, 0.01 pg/µL; G, 1 fg/µL; H, 0.1 f

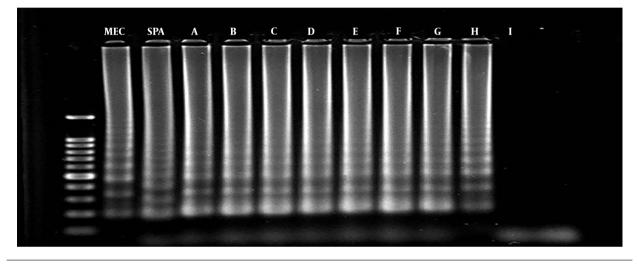


Figure 2. Multiplex LAMP products of spa and mecA genes. Tube 1, mecA positive control; tube 2, spa positive control; tubes 3 - 8, mecA and/or positive spaa specimens; tube 9, negative control.

4.3. Clinical Sensitivity and Specificity of Multiplex PCR and Multiplex LAMP

In this study, 53 clinical *S. aureus* suspected isolates were re-identified by standard biochemical tests, of which 49 isolates were *S. aureus* and four cases were non-aureus. According to the results of MIC of oxacillin sodium salt and antibiogram tests, 37 isolates were MRSA (oxacillin 4 > MIC μ g/mL) and 16 isolates (MIC < 2 μ g/mL) were MSSA. According to multiplex PCR, 49 isolates harbored *spa*, and the sensitivity and specificity were 100% compared to culture as the Gold standard.

Of the 49 isolates, 40 isolates harbored the *mecA* gene and showed as MRSA. Compared to the results of MIC and disk diffusion as Gold standards with a 95% confidence interval, the sensitivity, specificity, positive predictive value, and negative predictive value of multiplex PCR were 97.3%, 75%, 90%, and 31.92%, respectively. In multiplex LAMP, the sensitivity, specificity, positive predictive value, and negative predictive value of the test for the assessment of the *spa* gene were 100%, 100%, 100%, and 100%, respectively. Similarly, the sensitivity, specificity, positive predictive value, and negative predictive value of the test for the assessment of the *mecA* gene were 100%, 75%, 90.2%, and 100%, respectively.

4.4. Comparison of Multiplex LAMP and Multiplex PCR in Clinical Samples

As many as 40 strains harboring the *mecA* gene were observed in both methods, while 12 strains did not have the gene. Multiplex LAMP detected the *mecA* gene in one *mecA* multiplex PCR-negative isolate. Altogether, the sensitivity, specificity, positive predictive value, and negative predictive value of LAMP compared to PCR were 97.96%, 100%, 100%, and 92.31%, respectively. The kappa coefficient was calculated as 0.948.

5. Discussion

Methicillin-resistant *S. aureus* is a human pathogen that acts as both community and hospital-acquired infections worldwide. The increased MRSA frequency among health care centers has become a major problem in public health and infection control programs (14). Therefore, the rapid and accurate diagnosis of MRSA helps prevent the spread of nosocomial infections (25). Various methods like culture, PCR, MIC determination, and antibiogram are applicable for the detection of MRSA, although all of the mentioned methods suffer from some drawbacks. Molecular methods like PCR and real-time PCR are considered as methods of choice, as they are sensitive, accurate, and less time-consuming in comparison with culture and biochemical methods (26, 27).

Loop-mediated isothermal amplification is used for the rapid detection of DNA and RNA molecules. This technique, unlike PCR, does not need post-amplification procedures to detect the product and is advantageous compared to real-time PCR as it does not need expensive instruments (26, 27).

According to our experiment, the multiplex LAMP could detect the sequences of staphylococcal *mecA* and *spa* genes significantly more accurately than did multiplex PCR. This deduction was quite predictable considering similar publications, although the difference is promising. The calculated sensitivity of the multiplex LAMP in our study was much higher than that reported in similar experiments by Tie et al. and Su et al. (4, 17, 20). In a similar study conducted on the species of *Helicobacter*, higher percentages were reported for the sensitivity and specificity of multiplex LAMP than those of other methods such as multiplex PCR (21, 22).

Some of the features of the LAMP method include higher specificity and sensitivity and less time, labor, and equipment consuming than PCR. Moreover, since a large quantity of products is produced in the amplification by LAMP, the final amplicon detection can be performed without the need for gel electrophoresis equipment and it also allows for easy visualization by the naked eye (28). Typically, six primers (three pairs of primers) are used for the detection of about 100 bp of target DNA while, in PCR, two primers are attached to 40 bp. In this study, LAMP could detect the genes in clinical specimens in 188 minutes.

5.1. Conclusions

We introduced the multiplex loop-mediated isothermal Amplification as an alternative method for the diagnosis of methicillin-resistant *S. aureus*. This method has several advantages over PCR and culture, consisting of higher sensitivity and specificity. The authors would like to thank the staff of the Department of Microbiology, Miss Adabagher, and Mr. Safarian for their assistance.

Footnotes

Authors' Contribution: Ramin Abiri did study design and supervision. Amir Houshang Alvandi did primer design, supervision on LAMP. Sara Kashani did operation of the experiment, and writing of the manuscript.

Conflict of Interests: The authors declare no conflict of interest.

Ethical Approval: All procedures conducted in this study involving human participants and materials were approved by the Ethics Committee in Research, the Vice-Chancellor of Research and Technology, Kermanshah University of Medical Sciences (no.: 3005084)

Funding/Support: This project was supported by the Vice-Chancellor for Research and Technology, Kermanshah University of Medical Sciences (project no.: 92244).

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