A pooled analysis of the LAMP assay for the detection of Neisseria meningitidis

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Abstract

Background: Neisseria meningitidis is a major cause of bacterial meningitis, and these infections are associated with a high mortality rate. Rapid and reliable diagnosis of bacterial meningitis is critical in clinical practice. However, this disease often occurs in economically depressed areas, so an inexpensive, easy to use, and accurate technology is needed. We performed a pooled-analysis to assess the potential of the recently developed loop-mediated isothermal amplification (LAMP) assay for detection of meningococcus.

Methods: Pubmed, Embase, and Web of Science were searched to identify original studies that used the LAMP assay to detect meningococcus. After pooling of data, the sensitivity and specificity were calculated, a summary receiver operating characteristic (SROC) curve was determined, and the area under the SROC curve was computed to determine diagnostic accuracy. Publication bias was assessed using Deek’s funnel plot.

Results: We examined 14 studies within 6 publications. The LAMP assay had high sensitivity (94%) and specificity (100%) in the detection of meningococcus in all studies. The area under the SROC curve (0.980) indicated high overall accuracy of the LAMP assay. There was no evidence of publication bias.

Discussion: The LAMP assay has accuracy comparable to bacterial culture and PCR for detection of meningococcus, but is less expensive and easier to use. We suggest the adoption of the LAMP assay to detect meningococcus, especially in economically depressed areas.

Keywords: Neisseria meningitidis, Meningitis, LAMP assay, Pooled-analysis

Background

Neisseria meningitidis is a Gram-negative diplococci bacterium that only infects humans, and is a significant cause of meningitis. Neisseria meningitidis parasitizes in patient’s cerebrospinal fluid easily in the acute and early stage of diseases. But it is unknown of the carriage rate in healthy individuals at present [1]. Based on capsular polysaccharides, there are 13 serogroups of this species (A, B, C, D, E, H, I, K, L, W, X, Y, and Z) [2]. Serogroups A, B, C, Y, and W-135 are mainly responsible for human diseases, [3] and serogroups A and C are mainly responsible for meningitis. By the way we found other groups such as group B, Y and W135 cause severe epidemic lots of lands. The incidence of bacterial meningitis is greatest among infants under 1 year-old, followed by adolescent [4].

The overall incidence of meningitis has declined in recent decades due to the increasing use of meningococcal vaccines, [5] but the incidence of meningitis from bacterial strains not covered by vaccines has increased, and bacterial meningitis remains a significant public health problem. About 200,000 patients worldwide die from bacterial meningitis every year, and the mortality can be up to 60% in parts of sub-Saharan Africa and in poor
and developing countries [6, 7]. Patients who receive treatment have a mortality rate of about 10%, but survivors often experience serious sequelae, including limb amputation, neurological deficits, and other serious disabilities [8).

Clinical detection of Neisseria meningitidis in many countries currently uses PCR, ELISA, and bacterial culture [9] and PCR is considered the most authoritative standard. However, these technologies have limited applicability in poor and developing countries. For example, PCR and ELISA can be time-consuming, difficult to perform, and relatively expensive [10]. Identification by bacterial culture can take many days, has low sensitivity, and has a reduced detection rate for patients who were pretreated with antibiotics [6].

Loop-mediated isothermal amplification (LAMP) is a relatively new nucleic acid amplification technology that can simultaneously detect six meningococcal groups of N. meningitidis (A, B, C, W, X, and Y). It is simple to perform, rapid, has high sensitivity, and is inexpensive, [11] making it especially suitable for developing countries and regions with limited resources. No previous studies have comprehensively analyzed the sensitivity and specificity of the LAMP assay for detection of Neisseria meningitidis. The present analysis evaluated the diagnostic accuracy and feasibility of using LAMP to identify meningitis due to Neisseria meningitidis. This analysis also included the assessment of publication bias and data quality.

Methods
Study design
A systematic review of the diagnostic accuracy of LAMP in Neisseria meningitidis was performed, followed by a pooled-analysis.

Search strategy and study selection
The phrases “Neisseria meningitidis”, “Meningococcus”, “LAMP” and “Loop-mediated isothermal amplification” were used in combination to systematically search the literature from January 1997 to February 7, 2019 in 5 databases (Pubmed, Embase, Medline, Web of Science, and Cochrane Library) for identification of original studies that used LAMP to detect Neisseria meningitidis. The bibliographies of all publications were also reviewed to identify additional studies.

All literature titles and abstracts were independently screened, and five researchers (including Fan Shujin and Tan Hongkun) read the full text of each article that had extractable data to determine its eligibility for inclusion. Differences regarding eligibility were resolved through negotiation. Each included study used LAMP for detection of Neisseria meningitidis and a gold standard test (PCR or culture); provided data that were extractable and included true positivity (TP), false positivity (FP), false negativity (FN), and true negativity (TN) of the LAMP assay; and was original research written in English or Chinese. Studies that were animal experiments, from conferences or literature reviews, or did not have extractable data were excluded.

Data extraction and quality assessment
Five researchers independently extracted the data from each included study, including author, publication year, sample size, sample type, gold standard test results, TP, FP, FN, and TN. Regarding patient selection, index test, reference standard, flow and timing were used to evaluate the diagnostic accuracy of the LAMP assay.

Statistical analysis
MetaDisc version 1.4.0.0 was used to calculate sensitivity, specificity, positive likelihood ratio (PLR), negative likelihood ratio (NLR), and diagnostic odds ratio (DOR). Diagnostic odds ratio was used as the main outcome measure of the SROC approach of the standard method for analyzing diagnostic studies reporting pairs of sensitivity and specificity. A summary receiver operating characteristic (SROC) curve was plotted for calculations of sensitivity and specificity. A random effects model was used to summarize all data. Deek’s funnel plot was drawn using STATA, and a bivariate box plot and a post-test probability plot were used to detect literature bias. Review Manager version 5.2 was used for plotting study quality evaluation.

Results
Search results and study characteristics
Our initial literature search identified 83 publications and no additional (gray) literature (Fig. S1). After application of the inclusion and exclusion criteria, we identified 6 publications for inclusion in the analysis, all of which were published between 2011 and 2018 (Table 1). These studies were performed in Ireland, the United Kingdom, South Korea, and Japan. Because some of these publications used multiple types of sampling, we categorized these 6 publications as 14 studies with 3185 samples.

Diagnostic accuracy of LAMP assay
We used a random effects model to determine the diagnostic accuracy of the LAMP assay. Overall, the sensitivity was 0.94 (95% CI = 0.89 to 0.98, I² = 24.7%; Fig. 1), the specificity was 1.00 (95% CI = 0.99 to 1.00, I² = 58%; Fig. 2), the PLR was 69.45 (95% CI = 30.65 to 157.37, I² = 66.8%; Fig. 3), the NLR was 0.13 (95% CI = 0.08 to 0.21, and I² = 0.0%; Fig. 4), the DOR was 929.06 (95% CI = 339.22 to 2544.53, I² = 18.9%; Fig. 5), and SROC...
analysis indicated the Area Under Curve (AUC) was 0.9805 (Fig. 6).

**Quality evaluation**

We assessed the quality of the studies by analysis of patient selection, the index test, the reference standard, and study flow and timing (Figs. 7 and 8). The results indicated that patient selection had a high risk of bias in 3 studies, an unclear risk of bias in 3 studies, and a low risk of bias in 8 studies. Analysis of patient flow and timing indicated that only 1 study had a high risk of bias and 2 studies had an unclear risk of bias. Analysis of the index test indicated an unclear risk of bias in 1 study, and low risk of bias in the other 13 studies. Analysis of

**Table 1** Characteristics of the included studies

| Author          | Year | Country | Patients (n) | Specimen       | Reference standard | TP   | FP   | FN   | TN   |
|-----------------|------|---------|--------------|-----------------|-------------------|------|------|------|------|
| McKenna et al.  | 2011 | UK      | 139          | Throat swab     | PCR               | 12   | 1    | 0    | 126  |
| McKenna et al.  | 2011 | UK      | 141          | Serum           | PCR               | 14   | 1    | 0    | 126  |
| McKenna et al.  | 2011 | UK      | 72           | Blood           | PCR               | 4    | 1    | 0    | 67   |
| McKenna et al.  | 2011 | UK      | 22           | CSF             | PCR               | 3    | 1    | 0    | 18   |
| McKenna et al.  | 2011 | UK      | 16           | Respiratory secretions | PCR | 2    | 0    | 0    | 14   |
| McKenna et al.  | 2011 | UK      | 4            | Feces           | PCR               | 1    | 0    | 0    | 3    |
| Bourke et al.   | 2015 | UK      | 141          | Nasopharyngeal | PCR               | 21   | 0    | 4    | 116  |
| Bourke et al.   | 2015 | UK      | 144          | Blood           | PCR               | 22   | 0    | 4    | 118  |
| Lee et al.      | 2015 | South Korea | 31          | CSF             | Culture           | 2    | 3    | 0    | 26   |
| Lee et al.      | 2015 | South Korea | 568         | CSF             | PCR               | 15   | 5    | 0    | 548  |
| Lee et al.      | 2015 | South Korea | 536         | CSF             | PCR               | 5    | 0    | 0    | 531  |
| Lee et al.      | 2015 | South Korea | 470         | CSF             | PCR               | 5    | 1    | 0    | 464  |
| Takahashi et al.| 2016 | Japan    | 836          | Throat swab     | PCR               | 6    | 1    | 0    | 829  |
| Higgins et al.  | 2018 | UK       | 65           | Blood           | PCR               | 22   | 0    | 0    | 43   |

CSF cerebrospinal fluid, FN false negative, FP false positive, TN true negative, TP true positive

![Fig. 1 Sensitivity of the included studies](image-url)
Fig. 2 Specificity of the included studies

Fig. 3 Positive LR of the included studies
**Negative LR (95% CI)**

| Study     | LR      | 95% CI    |
|-----------|---------|-----------|
| McKenna   | 0.04    | (0.00 - 0.59) |
| McKenna   | 0.03    | (0.00 - 0.51) |
| McKenna   | 0.10    | (0.01 - 1.42) |
| McKenna   | 0.14    | (0.01 - 1.81) |
| McKenna   | 0.17    | (0.01 - 2.17) |
| McKenna   | 0.29    | (0.03 - 3.24) |
| Bourke    | 0.17    | (0.08 - 0.40) |
| Bourke    | 0.17    | (0.07 - 0.39) |
| Lee       | 0.19    | (0.01 - 2.38) |
| Lee       | 0.03    | (0.00 - 0.48) |
| Lee       | 0.08    | (0.01 - 1.19) |
| Lee       | 0.08    | (0.01 - 1.19) |
| Takahashi | 0.07    | (0.00 - 1.03) |
| Higgins   | 0.02    | (0.00 - 0.34) |

Random Effects Model
Pooled Negative LR = 0.13 (0.08 to 0.21)
Cochran-Q = 7.62; df = 13 (p = 0.8673)
Inconsistency (I-square) = 0.0 %
Tau-squared = 0.0000

**Diagnostic OR (95% CI)**

| Study     | OR      | 95% CI    |
|-----------|---------|-----------|
| McKenna   | 2,108.33 | (81.51 - 54,534.90) |
| McKenna   | 2,445.67 | (95.17 - 62,849.24) |
| McKenna   | 405.00   | (14.35 - 11,426.91) |
| McKenna   | 86.33    | (2.89 - 2,582.78)   |
| McKenna   | 145.00   | (2.30 - 9,137.08)   |
| McKenna   | 21.00    | (0.27 - 1,646.21)   |
| Bourke    | 1,113.22 | (57.82 - 21,434.61) |
| Bourke    | 1,185.00 | (61.63 - 22,783.90) |
| Lee       | 37.86    | (1.49 - 961.35)     |
| Lee       | 3,091.55 | (163.64 - 58,406.20) |
| Lee       | 11,693.00| (212.26 - 644,149.89) |
| Lee       | 3,406.33 | (124.49 - 93,206.44) |
| Takahashi | 7,189.00 | (267.19 - 193,427.75) |
| Higgins   | 3,915.00 | (75.17 - 203,905.95) |

Random Effects Model
Pooled Diagnostic Odds Ratio = 929.06 (339.22 to 2544.53)
Cochran-Q = 16.04; df = 13 (p = 0.2472)
Inconsistency (I-square) = 18.9 %
Tau-squared = 0.6969
reference standards indicated an uncertain risk of bias in 4 studies and low risk of bias in 9 studies. Analysis of applicability concerns indicated low concerns in all three categories.

**Publication bias**
Deek's funnel plot (Fig. 9) showed that publication bias was not significant ($P = 0.07$). A bivariate box plot of sensitivity and specificity showed that there was no significant heterogeneity among the included studies (Fig. 10). The post-test probability nomogram showed that for samples with a predicted probability of 50%, the post-test probability of a positive result was 100%, and the post-test probability of a negative result was 1% (Fig. 11).

**Discussion**
Bacterial meningitis caused by *Neisseria meningitidis* has high global incidence and mortality rates, and is an especially severe problem in undeveloped regions. At present, routine laboratory diagnosis of bacterial meningitis employs bacterial culture and Gram staining, but this method is limited because it has a low sensitivity and is very time-
consuming [18, 19]. PCR is a widely accepted laboratory test for diagnosis of bacterial meningitis, and many studies have confirmed its high sensitivity and specificity. Unfortunately, the current PCR applications used to detect bacterial meningitis are too complicated and expensive for clinicians in undeveloped regions [19, 20]. Thus, existing PCR tests are difficult to apply in areas with the highest incidences of bacterial meningitis. A rapid, low-cost, easy to use, and highly sensitive technique is needed for these at-risk populations.

LAMP is a simple technique developed in recent years that uses 4 specific primers in 6 regions of the target gene and does not require a thermocycler. Under the action of a strand displacement DNA polymerase (Bst DNA polymerase), genes are amplified at a constant temperature of 60 to 65 °C, and amplification can be 10⁹-fold to 10¹⁰-fold in 15 to 60 min. Higgins et al. [17] used Tth endonuclease IV and a unique LAMP primer/probe to develop novel real-time multiplex LAMP technology, TEC-LAMP. Its main advantages are that it is simple, fast, and inexpensive [12, 21].

We performed a pooled-analysis of 14 independent studies reported in 6 publications. These studies were performed in diverse geographic regions and used a variety of different samples. Thus, the study of McKenna et al. [13] was from Ireland and had 139 throat swab samples, 141 plasma samples, 72 blood samples, 22 cerebrospinal fluid samples, 16 respiratory secretion samples, and 4 stool samples. The study of Bourke et al. [14] was from the United Kingdom and had 141 nasopharyngeal swab samples and 144 blood samples. The study of Lee et al. [15] was from South Korea and examined 31 cerebrospinal fluid samples. The study of Lee et al. [16] was also from South Korea and examined 1574 cerebrospinal fluid samples (568 from Vietnam, 536 from China, and 470 from

![Fig. 8 Quality assessment of the individual studies](image-url)
South Korea). The study of Takahashi et al. [1] was from Japan and examined 836 throat swab samples. The study of Higgins et al. [17] was from Ireland and examined 65 blood samples. The study of Lee et al. [14] used culturing as the reference standard, and all the other studies used PCR as the reference standard. Our analysis results indicated that LAMP had an overall sensitivity of 94% (95% CI = 0.89 to 0.98) and an overall specificity of 100% (95% CI = 0.99 to 1.00) for the detection of *N. meningitidis*, and that the accuracy was high regardless of the sample type and geographic origin of the patients. Therefore, LAMP is highly suitable for the rapid, accurate, and inexpensive clinical detection of *N. meningitidis* in economically impoverished areas. Further prospective studies are needed to verify the safety and most suitable samples for the clinical use of LAMP.

**Conclusions**

In conclusion, our research showed that the LAMP assay had high sensitivity and specificity in detecting bacterial
meningitis due to *N. meningitidis*. This assay thus provides reliable information for clinical laboratory tests, and is suitable for detecting bacterial meningitis. However, further research is needed to verify the clinical feasibility and accuracy of this test in the impoverished areas where it is most needed.

**Supplementary information**

Supplementary information accompanies this paper at https://doi.org/10.1186/s12879-020-05250-w.

**Abbreviations**

LAMP: Loop-mediated isothermal amplification; SROC: Summary receiver operating characteristic; PCR: Polymerase Chain Reaction; ELISA: Enzyme Linked Immuno Sorbent Assay; TP: True positivity; FP: False positivity; FN: False negativity; TN: True negativity; PLR: Positive likelihood ratio; NLR: Negative likelihood ratio; DOR: Diagnostic odds ratio

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**Authors’ contributions**

XGG, FSJ, THK, XYC, CYZ, XTA, PZY, OS, LQ, LXY and LZX participated in the design of the project, formulation of the search strategy, and determination of inclusion of exclusion criteria. All authors participated in the literature search, data extraction and processing, and quality evaluation. XGG has made substantial contributions to the conception and design of the work.
FSJ created the figures and table. FSJ and THK wrote the manuscript. All researchers read and approved the final version of the manuscript.

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Availability of data and materials
All data generated or analyzed during this study are included in this published article and its supplementary information files.

Ethics approval and consent to participate
Not applicable.

Consent for publication
Not applicable.

Competing interests
There is no competing interests in research.

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