

# Clinical Value of SARS-CoV2 IgM and IgG Antibodies in Diagnosis of COVID-19 in Suspected Cases

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**Objective:** To explore the clinical value of SARS-CoV2 IgM and IgG antibodies in the diagnosis of COVID-19 in suspected cases by likelihood ratio.

**Methods:** By reinterpreting data from a previous study, the positive likelihood ratio of IgM and IgG antibodies in COVID-19 pneumonia diagnosis was calculated, and the posterior probability of IgM and IgG antibodies and their tandem detection was calculated finally.

**Results:** The positive likelihood ratios of single IgM and IgG antibodies were 18.50 and 12.65, respectively, and the posterior probabilities were 90.18% and 86.26%, respectively. However, the posterior probability of the two antibody-tandem test was 99.15%, which could give clinicians more quantitative confidence in the diagnosis of COVID-19 in suspected cases. According to the results of this study, combining the advantages and disadvantages of nucleic acid testing and antibody detection, a feasible clinical path was found for clinicians to diagnose COVID-19 pneumonia from suspected cases.

**Conclusion:** For suspected cases, IgM- and IgG-antibody tests should first be done at the same time. If all antibody tests are positive, COVID-19 pneumonia could be confirmed. If not, nucleic acid detection (once or more) should be carried out, and in extreme cases high-throughput viral genome sequencing is required.

**Keywords:** COVID-19, SARS-CoV2, positive likelihood ratio, posterior probability, antibody test, nucleic acid test, clinical pathway

## Introduction

Since the outbreak of SARS-CoV2 last December in China, >51 million cases have been confirmed in >180 countries. As the country with the most severe epidemic in the early period, China has accumulated a lot of experience in the diagnosis and treatment of COVID-19.<sup>1,2</sup>

Based on increased awareness of the disease, the treatment plan for COVID-19 has been constantly updated by the Chinese government. The latest two versions<sup>3,4</sup> are versions 7 and 8. In version 7, the standard indicators used for confirmed diagnosis of suspected cases were nucleic acid detection, gene sequencing, and antibody testing for SARS-CoV2. Of these, antibody testing was first among the diagnostic criteria. In version 8, antibody testing as a diagnostic indicator of suspected cases was further confirmed. On April 2, 2020, the US Food and Drug Administration approved the first SARS-CoV2 antibody-test kit for COVID-19 detection. The Chinese National Medical Products Administration had also approved 25 antibody-test kits by November 11. Test methods included colloidal gold immunochromatography and chemiluminescence.

Bayesian analysis is based on the sensitivity and specificity of diagnostic testing, combined with the proportion of various diseases in the population (prior probability)

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and calculating the probability of suffering from various diseases (posterior probability). Posterior probability can provide a basis for individual diagnosis, which could help clinicians in interpreting test results scientifically and improve diagnosis.<sup>5</sup> Posterior probability in antibody testing to diagnose COVID-19 in suspected cases has not been reported, so this study intended to use likelihood ratios (LRs) with antibody testing to calculate posterior probability to clarify the significance of antibody testing in the diagnosis of suspected cases.

In addition, how to use nucleic acid detection and antibody test to diagnose suspected COVID-19 patients more efficiently was a question worth pondering. As such, this study also intended to propose a feasible clinical pathway for clinicians to diagnose suspected cases by nucleic acid and antibody detection.

## Methods

### Data Sources

The data in this study come from articles on COVID-19 published online by the *Chinese Journal of Laboratory Medicine*, mainly Ming et al<sup>6</sup> and Wanzhou et al.<sup>7</sup> The subjects were all suspected and confirmed cases of COVID-19 admitted to the People's Hospital Affiliated with Wuhan University.

### Case Definitions

#### Suspected Cases: Epidemiological History and Clinical Manifestations

Epidemiological history comprises history of travel to or residence in Wuhan and surrounding areas or other communities where cases have been reported within 14 days prior to the onset of the disease, contact with novel coronavirus-infected people (with positive results on nucleic acid testing) within 14 days prior to the onset of the disease, in contact with patients who have fever or respiratory symptoms from Wuhan and surrounding area or from communities where confirmed cases have been reported within 14 days before the onset of the disease, and clustered cases (two or more cases with fever and/or respiratory symptoms in a small area, such as families, offices, and schools, within 2 weeks.

Clinical manifestations comprise fever and/or respiratory symptoms, the aforementioned imaging characteristics of COVID-19, normal or decreased white blood-cell count, and normal or decreased lymphocyte count in the early stage of onset. A suspect case has any of the epidemiological history plus any two clinical manifestations or

all three clinical manifestations if there is no clear epidemiological history.

### Confirmed Cases

Suspect cases are those one of the following etiological or serological findings: real-time fluorescent PCR positive for SARS-CoV2 with nucleic acid, Viral gene sequence highly homologous with known SARS-CoV2, SARS-CoV2-specific IgM and IgG detectable in serum, and SARS-CoV2-specific IgG detectable or reaches a titration of at least fourfold the acute phase during convalescence.

### Data and Testing

Positive LR (LR<sup>+</sup>) of IgM and IgG in suspected cases for nucleic acid-positive patients was calculated mainly by reinterpreting the data from Wanzhou et al.<sup>7</sup> According to relevant data from Ming et al,<sup>6</sup> the posterior probability of IgM- and IgG-antibody detection alone and in series for diagnosis of nucleic acid-positive patients was calculated. SARS-CoV2 IgM and IgG were tested with chemiluminescence immunoassay, and nucleic acid tested using RT-PCR.

### Statistical Methods

Data were calculated using a four-grid table.

## Results

### Diagnostic Performance of IgG and IgM in Diagnosing COVID-19

Wanzhou et al<sup>7</sup> found that 205 patients had coronavirus infection and 79 patients no COVID-19 pneumonia. Combined with the number of IgG- and IgM-antibody detection mentioned in this article, [Tables 1](#) and [2](#) were obtained.

As shown in [Table 1](#), sensitivity, specificity, negative predictive value, and positive predictive value of SARS-CoV2-antibody detection in the diagnosis of pneumonia in COVID-19 were 96.10% (197 of 205), 92.40% (73 of 79), 90.10% (73 of 81) and 97.04% (197 of 203), respectively.

**Table 1** Diagnostic Performance of IgG Testing to Diagnose COVID-19

	COVID-19		Total
	Positive	Negative	
Positive	197	6	203
Negative	8	73	81
Total	205	79	284

**Table 2** Diagnostic Performance of IgM Testing to Diagnose COVID-19

	COVID-19		Total
	Positive	Negative	
Positive	144	3	147
Negative	61	76	137
Total	205	79	284

The  $LR^-$  was 0.04 and the  $LR^+$  12.65. Accuracy was 95.07% (270 of 284).

As shown in Table 2, sensitivity, specificity, negative predictive value, and positive predictive value of SARS-CoV2 IgM-antibody detection to diagnose COVID-19 were 70.24% (144 of 205), 96.20% (76 of 79), 56.72% (76 of 137), and 97.76% (144 of 147), respectively. The  $LR^-$  was 0.31 and the  $LR^+$  18.50. Accuracy was 77.46% (220 of 284).

## Posterior Probability of Single-IgG and -IgM and IgM-IgG Tandem Diagnostic Tests

For posterior probability of a single index:

$$\text{prior ratio} = \text{prior probability}/(1 - \text{prior probability})$$

$$\text{posterior ratio} = \text{prior ratio} \times LR^+$$

$$\text{Posterior probability} = \text{posterior ratio}/(1 + \text{posterior ratio})$$

For posterior probability formula of tandem diagnostic tests of multiple indices:

$$\text{prior ratio} = \text{prior probability}/(1 - \text{prior probability})$$

$$\text{posterior ratio} = \text{prior ratio} \times LR^+ 1 \times LR^+ 2 \times \dots \times LR^+ n$$

$$\text{Posterior probability} = \text{posterior ratio}/(1 + \text{posterior ratio})$$

According to the 8,274 suspected cases of COVID-19 explicitly mentioned in Ming et al,<sup>6</sup> the proportion of confirmed cases from suspected cases was 33.17%, which was the prior probability. The posterior probability of single testing of IgG and IgM and IgM-IgG tandem diagnostic testing is shown in Table 3.

**Table 3** Posterior Probability of Diagnostic COVID-19 on Single-IgG and -IgM and IgM-IgG Tandem Tests

	Prior Probability	Prior Ratio	Posterior Ratio	Posterior Probability
IgG	33.17%	0.4963	6.28	86.26%
IgM	33.17%	0.4963	9.18	90.18%
IgM-IgG	33.17%	0.4963	116.19	99.15%

For suspected cases, after single detection of IgG and IgM antibodies, the clinician's confidence in the diagnosis of suspected patients as confirmed COVID-19 cases was 86.26%– 90.18%, but after IgM-IgG tandem testing, the clinician's confidence in the diagnosis of suspected patients as confirmed COVID-19 cases increased to 99.15%.

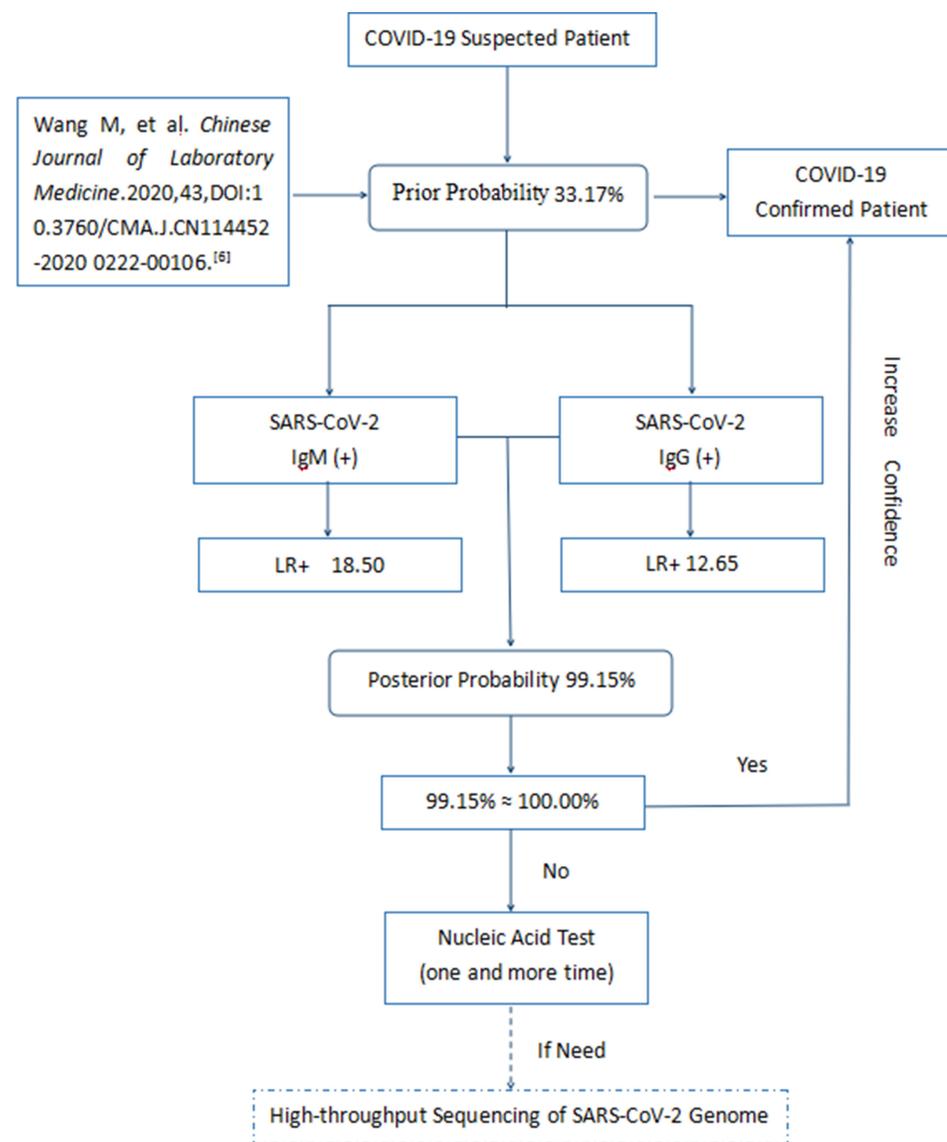
## Clinical Pathway for Clinicians to Diagnose COVID-19 Based on IgM-IgG Antibody-Tandem Testing

According to the results of this study, combining the advantages and disadvantages of nucleic acid detection and antibody detection, the clinical pathway for clinicians to diagnose COVID-19 is shown in Figure 1. It puts forward a feasible clinical pathway for the diagnosis of pneumonia in COVID-19, starting with IgM-IgG tandem test, then using nucleic acid detection, and finally carrying out high-throughput sequencing if necessary.

## Discussion

There is continuous development detection methods for human antibodies after SARS-CoV2 infection.<sup>8–10</sup> Nucleic acid and antibody detection are the main laboratory testing methods for COVID-19, but have their own advantages and disadvantages.<sup>11,12</sup> First of all, from the aspect of specimen sampling, commonly used specimens for nucleic acid detection are sputum, nasopharynx swab, or various lung-lavage fluids. Sampling poses great infection risk to medical personnel and also potential infection risks for laboratory personnel for nucleic acid detection.<sup>13</sup> However, if the type of antibody-test specimen is blood, which is relatively easy to obtain, the risk of infection of medical personnel is relatively low. Secondly, according to the quality of the detection results, nucleic acid detection is restricted by the quality of sample materials and the extraction quality of nucleic acid, and there can be serious undetected phenomena. Different sample types have different detection rates.

Bronchoalveolar lavage-fluid specimens show the highest positive rates (93%), followed by sputum (72%), nasal swabs (63%), fibrobronchoscope brush biopsy (46%), pharyngeal swabs (32%), feces (29%), and blood (1%).<sup>14</sup> Repeated nucleic acid detection for some suspected patients is necessary and can effectively avoid false-negative result. For antibody detection, except for some patients with low immunofunction, antibodies cannot be produced effectively. Antibodies should be present in most infected persons.<sup>15</sup> Antibody detection is limited



**Figure 1** Clinical pathway for clinicians to diagnose COVID-19 by SARS-CoV2 antibody.

mainly by the sensitivity and related performance of the kit. In addition, there will be some false positives caused by detection interference in tumor patients, patients with autoimmune diseases, and others with chronic infection. Finally, the convenience of antibody detection has advantages that nucleic acid detection does not have at all, including simplicity and convenience of operation, qualifications of personnel, and requirements of the detection environment. Therefore, it would be of great clinical value to study how to optimize antibody detection and nucleic acid detection in clinical diagnosis.

No clear research on feasible methods of antibody or nucleic acid detection has been found yet. Therefore, this research used published data and adopted the method of

$LR^+$  to calculate posterior probability, in order to propose the possibility of optimizing the diagnosis pathway for the first time. Since the original data of the two-antibody tandem test could not be obtained from Wanzhou et al,<sup>7</sup> the diagnostic performance index of the tandem test could not be analyzed in this paper. However, this study solved that problem by calculating posterior probability through the  $LR^+$ , and proved that the posterior probability of the two-antibody tandem test was much higher than each single-antibody detection; however, the posterior probability of single-antibody detection was not different. This showed that the single-antibody test could not significantly increase the confidence of clinicians in a clear diagnosis of confirmed cases from suspected cases. However, the two-

antibody tandem test obviously improved this confidence, raising clinicians' confidence in diagnosis to 99.15%, which was helpful in optimizing the diagnostic process through quantification. The clinical pathway for clinicians to diagnose COVID-19 for suspected cases should be firstly conducting IgM- and IgG-antibody tests. If all antibody tests are positive, COVID-19 can be confirmed. If not, nucleic acid detection (one or more times) can be performed, and in extreme cases high-throughput viral genome sequencing.<sup>16</sup>

In short, the evaluation of SARS-CoV2 IgM and IgG antibodies serves epidemiological purposes for COVID-19 that are recognized.<sup>17</sup> Many conventional diagnostic studies have reported the diagnostic significance of antibody testing,<sup>18–20</sup> but in the general population antibody test should not be used as a diagnostic index. It has diagnostic significance only for suspected cases. Compared with the traditional diagnostic performance index, the posterior probability calculated by LR<sup>+</sup> can better reflect serial detection of the two antibodies. In practice, it will be more helpful for clinicians in optimizing the diagnostic process. This study was aimed at the secondary excavation of published literature data. In future, the original data should be used and the sample size expanded to further verify the conclusion.

## Disclosure

The author declares no conflicts of interest.

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