

## Severe Acute Respiratory Syndrome -Possible Sources of Error in Diagnosis and their Validity in the Coronavirus Disease Series

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#### **Mini Review**

Volume 7 Issue 2 Received Date: June 06, 2022 Published Date: August 04, 2022 DOI: 10.23880/ijbp-16000212

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#### Abstract

Viral diseases (recently monkeypox viral disease) are very common in our era and some viruses such as SARS-CoV-2 threaten humanity by causing a severe acute respiratory syndrome. This disease has been linked to haematological dysfunctions. Spike (S) and Nucleoprotein (N) proteins have been putatively associated with these dysfunctions. There is no currently fully effective detection method and treatment. Therefore, this study highlighted coronavirus disease's diagnosis and prevention methods (COVID-19), an infectious disease caused by a severe acute respiratory syndrome virus. Because the coronavirus strain of severe acute respiratory syndrome has been threatening humans since 2003 and reappeared in 2019 at the end of 2019, causing many deaths by 2022 under the name coronavirus disease. Because the coronavirus disease rapidly spread across the world due to its high transmissibility and prolonged incubation. Therefore, it seems that SARS coronaviruses in the form of SARS-CoV-3, SARS-CoV-4, etc., will not disappear from our world in the future. Thus, in this study, possible sources of error in diagnosis and their validity that can be used today and in the future in the SARS coronavirus disease series are presented in the light of current information.

**Keywords:** Analytical Error; COVID-19 Disease; RT -PCR; SARS Coronavirus Disease

**Abbreviations:** S: Spike; N: Nucleoprotein; RT-PCR: Real-Time Reverse Transcription-Polymerase Chain Reaction; BAL: Bronchoalveolar Lavage Specimen; FB: Flexible Bronchoscopy.

#### Introduction

#### Potential Sources of Error in the Diagnosis of the SARS Coronavirus Disease Series and Validity of the Methods

The SARS coronavirus disease series (including the disease COVID-19) is continuously threatening humanity [1].

In order to get rid of the SARS coronavirus disease series, the correct diagnosis must first be made. However, previous studies have shown that the reliability and validity for such diseases are extremely low, except for real-time reverse transcription-polymerase chain reaction (RT-PCR) [2,3]. In this study, we aim to explore in detail the reasons that affect the validity and reliability of the assays. There are three sources of error in analytical laboratories: preanalytical, analytical, and post-analytical. For a test to be accurate and reliable, all three sources of error must be eliminated. Possible sources of error were given below one by one.

#### **Preanalytical Sources Of Error**

In laboratory results, most errors are made at this stage. The most important of these errors are: insufficient knowledge and experience of the personnel, improper preparation of the patient, wrong barcode, missing sample, wrong sample, inappropriate tube, clotted sample, hemolyzed or lipemic sample, contamination, samples are not delivered to the laboratory under appropriate conditions and in the appropriate time frame [4]. In the diagnosis of SARS coronavirus disease series (including COVID-19), the most likely preanalytical errors are errors in collecting nasopharyngeal swabs, throat swabs, and, in later stages of the disease, bronchoalveolar lavage specimens from the field. Because blood specimens are collected and analyzed continuously in all laboratories, no unusual errors are expected in collecting and analyzing these specimens, except for staff inexperience and carelessness [5-9].

However, because errors are more likely to occur in the collection of nasopharyngeal swabs and throat swabs for diagnosis of the SARS coronavirus disease series (including COVID-19), and bronchoalveolar lavage specimens in the later stages of the disease, these biological specimens should be used to collect biological specimens in this part of the study correctly [9-11].

#### **Collection of a Nasopharyngeal Swab**

To collect a nasopharyngeal swab, first cover the patient's mouth with a mask, leaving the nose open. The patient should then be asked which nostril they are breathing through. If your patient breathes through both nostrils, you can collect the specimen by choosing one of the two nostrils. The head should be extended when collecting the sample, and the sample stick advanced to the nasopharynx. When the sample stick touches the nasopharynx, the sample should be collected by slowly rotating the sample stick around itself after waiting for a few seconds to sufficiently absorb the secretion. After collecting the specimen, the portion containing the nasopharyngeal swab should be broken and placed in a sterile test tube. Personnel carrying the specimen should be sent to the laboratory immediately, providing information about the nature of the specimen and warning of contamination [12].

#### **Collection of a Throat Swab**

To collect a throat swab, the patient's mouth must first be covered with a mask to keep it open. The head is then stuck out, and after the head is stuck out, gentle pressure is applied to the tongue with a tongue depressor under an adequate light source. A sterile swab is then taken by moving the swab back and forth between the two tonsils behind the uvula without touching the lips, buccal mucosa, uvula, and tongue by moving the swab back and forth in the posterior pharynx and tonsil area for 5 seconds and should be sent to the laboratory [13-16].

# Collection of Bronchoalveolar Lavage Specimen (BAL)

This biological sample should be collected by an experienced physician using a bronchoscope. The patient whose collected bronchoalveolar lavage sample should not eat for at least 6 hours. Before taking the sample, a local anesthetic (prilocaine hydrochloride 2%; 5-10 mL) should be applied with the help of a nebulizer, and then a local anesthetic (lidocaine 2% spray) should be administered in the form of an oropharyngeal and oropharyngeal spray (to avoid the side effects of local anesthetics), the dose should not exceed 4 mg/kg. In some cases, mild sedation with benzodiazepines or with the aid of a morphine analog may be required. This is at the discretion of the physician. However, many clinicians recommend sedation before the procedure as it improves airway assessment and returns to BAL. Flexible bronchoscopy (FB) is initially inserted transnasally or transorally for this procedure. If there is involvement, BAL should be applied to the right middle lobe segment, partial segments, or the left upper lobe lingual segments. To the involved segment, FB mouthpiece (wedge position), 100-300 ml of 0.85% NaCl (saline) should be injected bronchoscopically into the canal in 50-ml portions and taken by hand via the injector. Care should be taken to ensure that the physiologic serum used for lavage is at body or room temperature. In addition, at least 70% of the administered fluid should be returned. To increase the return flow, too high pressure should not be applied. This is because high pressure leads to deterioration of the contents of the bronchoalveolar lavage material. The sample collected in this way reflects approximately one million alveoli. Because the agent of the SARS coronavirus disease series (including COVID-19) is sought in this biological material, it should not be filtered (in other cases where cellular contents are to be examined, it should be filtered through several layers of gauze to clear it of mucus and transferred to containers of

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ice to preserve cell viability). Samples from different alveoli and distal bronchioles can be collected in the same container. Samples should be collected at the time when the virus can be detected. Nasopharyngeal and pharyngeal swabs, saliva, and blood should be the first choice biological specimens for screening for this virus. BAL is a semi-invasive diagnostic method used to diagnose infectious and non-infectious diseases. The most important thing is to completely disinfect the bronchoscope and perform all procedures under proper safety standards when collecting BAL. 2-3 ml of the tracheal aspirate, bronchoscopic specimens thus obtained should be immediately stored in a sterile tube with a screw cap or an injector sealed with a Luer cap in the refrigerator (between 2-8 0 C) and sent to the laboratory as soon as possible, and should be searched for the cause of the SARS coronavirus disease series (including COVID-19). If desired, specimens may be stored at -20°C or -70°C if feasible. Samples stored at -20°C can remain intact with cell contents for two months. At -70°C, on the other hand, samples can be stored longer.

In addition to the biological samples mentioned above, the blood sample is an important fluid in diagnosing the SARS coronavirus disease series (including COVID-19). However, because this biological fluid is used worldwide, the margin of error is low. Therefore, no information has been provided on how to collect blood samples. If you have any doubts, also ask the laboratory personnel. When collecting these biological fluids that we use for diagnosis, there are always analytical errors in laboratories [10,17].

#### **Possible Analytical Errors**

The main possible causes of analytical errors are

- Uncalibrated equipment
- Bubbles or particles in the samples that are out of range
- Rrroneous deviations
- The presence of undefined molecules interferes with the method used or insufficient sensitivity of the kits used to measure specificity and sensitivity.

Elimination of these analytical errors will allow accurate results.

#### **Potential Post-Analytical Errors**

Reporting delays include forwarding results to another site or reporting results incorrectly. Methods Currently Used for Diagnosis

#### **Biochemical Parameters**

Currently, no biochemical parameters can be measured with fully sensitive auto analyzers. However, serum, urea, keratin, cystatin C, cholinesterase, direct bilirubin, lactate dehydrogenase, interleukin-6, serum ferritin, D-dimer, and coagulation cascade parameters appear to be among the biochemical parameters of the SARS coronavirus disease series (including COVID-19) [18-24].

#### **Chest CT Scan and Radiography**

Since SARS-CoV-2 causes fibrosis in the lungs of the SARS coronavirus disease series (including COVID-19), lung radiographs provide information about the status of the disease. It has been reported that these lung films, in conjunction with artificial intelligence programs, help diagnose COVID-19 with 60-70% accuracy [25,26].

#### **Diagnosis by Antibody Measurement**

Many different companies have produced very different kits for antibody measurement. However, no kit has been produced to date that definitively supports the diagnosis. The sensitivity of some kits is so low as to be almost negligible. This is probably due to the constant mutation of the SARS coronavirus series (including COVID-19), which leads to measurement uncertainties [27]. Currently, three different SARS-CoV-2, A, B, and C forms cause infections in different countries. Therefore, it seems almost impossible to measure a constantly mutating virus using the antigenantibody principle-based method, and it can only partially inform physicians [28]. The principles of this method and how it works are explained in a previous study [4].

#### **Diagnosis by Measurement Using RT -PCR**

The causative agent of the SARS coronavirus series (including COVID-19) is currently definitively identified by viral RNA detection using RT-PCR. However, it is important to remember that false-negative results are possible when biological samples are collected before the disease's incubation period. This method is sensitive enough to detect viral RNA in asymptomatic cases. Until viral PCR testing is confirmed, laboratories need to provide detection by pan-coronavirus testing followed by sequence analysis. It is important to exclude other coronaviruses that may prove positive upon confirmation, especially pancoronavirus tests (HCoV-229E, HCoV-NL63). This is because beta-coronaviruses, SARS, and MERS-CoV viruses cause zoonotic infections in humans [29,30]. The above methods are currently being used to diagnose COVID-19 disease. However, let us take a look at the scientific knowledge of our world so far. Other biochemical parameters may play a role in the possible diagnosis of the disease in the SARS coronavirus disease series (including COVID-19). Therefore, in this part of the study, we will look at the method and some biochemical parameters that can give us an idea of the SARS coronavirus disease series (including COVID-19) and could be applied in laboratories' future.

#### Other Possible Methods and Parameters that Could Aid in Diagnosis Include the Following

Hemagglutination Inhibition Test: Members of the Corona family of viruses can agglutinate erythrocytes. In a 1979 study of cattle, the presence of coronavirus was determined by the hemagglutination inhibition test. This test is very simple [31]. An alternative test is available in countries that cannot perform the aforementioned advanced tests. This test should be compared to existing sensitive tests for its sensitivity and adapted to human plasma and serum. All detailed information about the test can be found in the Wang, et al. study [32]. In brief, erythrocyte agglutination has been detected in patients who died because of COVID-19. Therefore, checking whether red cells are agglutinated may also help diagnose COVID-19 disease. To prevent erythrocyte agglutination, the administration of anti-erythrocyte agglutinators to these patients is also expected to reduce clot-related deaths by eliminating agglutination. In addition, it is predicted that the administration of low-weight heparin to these patients will also be beneficial [32].

Hemoglobin Electrophoresis Test: Because SARS-CoV-2, one of the SARS coronaviruses, now targets the beta chain of hemoglobin, hemoglobin electrophoresis may also aid in diagnosis [33]. Theoretically, a break in the beta chain shows a different band in hemoglobin electrophoresis [34-36]. However, evaluating hemoglobin electrophoresis and a complete blood count gives a better indication. Since the beta attack of SARS-CoV-2 hemoglobin is valuable information [37], we think it is useful to provide brief information on this topic. SARS-CoV-2 attacks the beta chain of hemoglobin, which is responsible for oxygen transport to tissues and cells by disrupting the structure of porphyrin and inhibiting heme synthesis. It has been reported that patients with deceased COVID-19 have more than twice as much serum ferritin in their circulation [38]. Iron deficiency and excess are pathological conditions. Therefore, iron metabolism must be strictly controlled. In mammals, iron balance is controlled at the level of intestinal absorption. Hepcidin, a peptide hormone, is the major regulator of systemic iron balance. Hepcidin maintains iron balance by coordinating the use and storage of iron and preventing its leakage into plasma. Hepcidin reduces iron absorption from the small intestine. Hepatic hepcidin synthesis decreases with increased erythropoietic activity, hypoxia, and decreased iron storage in the organism. On the other hand, hepcidin synthesis increases in the presence of iron overload or inflammation [39,40].

#### Other Possible Biochemical Parameters that May Play a Role in the Diagnosis

**Hepcidin:** Since SARS-CoV-2, one of the SARS coronavirus disease series, increases iron by binding to the beta portion of

hemoglobin, the amount of hepcidin will inevitably increase due to free iron [41,42]. When we combine this information, the measurement of hepcidin accumulation represents a new biomarker candidate indicating the presence of COVID-19. In addition, we are not in favor of suppressing the increased iron due to COVID-19, which we mentioned here. This is because the Fenton reaction forms free radicals in large quantities [43]. Although free radicals are harmful to the organism, they also have a benefit due to their bacteriostatic effect [43,44]. The increased iron in COVID-19 associated with the disease may play an important role in killing bacteria via the enzyme myeloperoxidase [45], so we believe suppressing the increased iron in the circulation by COVID-19 is not the right option.

**Maresin-1:** Maresin 1, this molecule, which is a lipid regulator, also has anti-inflammatory properties. In studies of the SARS coronavirus disease series, inflammation levels have been elevated, especially at COVID-19. Therefore, it is thought that Maresin 1 may be a useful biomarker for diagnosing this disease [46,47].

**Relaxin:** Since relaxin is a molecule that decreases inflammation [48,49], we think it is useful to measure this molecule in COVID-19 when inflammation is elevated.

**Measurement of Surfactants:** Patients with COVID-19 experience respiratory failure syndrome. This could be due to a decrease in surfactants in the lungs' alveoli [50]. Therefore, it is predicted that the measurement of surfactants in COVID-19 may aid in diagnosis. In addition, it is predicted that the replacement of surfactant will help keep the alveoli open by decreasing surface tension and may also eliminate respiratory failure syndrome. If the SARS coronavirus disease series occur in the future, measurement of surfactants in these series may also be beneficial.

#### Conclusion

If the SARS coronavirus disease series continues in the future, in that case, control of the disease will play an important role in the rapid and accurate diagnosis and treatment of this disease, as well as following the recommendations applied by countries and recommended to their citizens. However, as mentioned above, the most important role in controlling the disease depends on the correct diagnosis of the laboratories and the treatment protocols followed by physicians depending on this diagnosis. It is believed that eliminating the potential sources of error identified in this study will help ensure that the diagnosis is made quickly and accurately. This will be a means of quickly bringing the epidemic under control.

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