

Role of the Laboratory in the Outbreak, Spread, Prevention and Control of Covid-19 Pandemic in Nigeria

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Abstract

Coronavirus disease 2019, abbreviated to COVID-19, is the latest biological hazard to assume the relevance of insidious worldwide threat. The responsible pathogen is a virus belonging to the Coronaviridae family, finally defined as “severe acute respiratory syndrome coronavirus 2” (SARS-CoV-2) for high sequence identity (i.e. up to 80%) with the homologous virus which caused the SARS outbreak in 2003 (i.e. SARS-CoV). At the time of writing this article, SARS-CoV-2 has already infected over 115,000 people in more than 115 different countries, causing nearly 4000 related deaths. Structural analysis shows that SARS-CoV-2 probably derives from a bat SARS-like coronavirus, which has been then transmitted to humans after emergence of mutations in the spike glycoprotein (protein S) and nucleocapsid N protein. The mutation that occurred in the former protein is especially important, whereby viral spike glycoprotein mediates the entrance of the virus into the cell through cell receptor binding and membrane fusion. On the other hand, the N protein regulates the process of viral replication, thus influencing transcription and assembly. Altogether, mutations in these two proteins would then explain the unique characteristics of SARS-CoV-2 compared to the original SARS-CoV, i.e. enhanced infectious potency in humans, combined with relatively mitigated pathogenicity.

Keywords: *Coronaviridae; Nucleocapsid; Pathogenicity; Glycoproteins; Biowarfare*

Introduction

Coronavirus disease 2019, abbreviated to COVID-19, is the latest biological hazard to assume the relevance of insidious worldwide threat. The responsible pathogen is a virus belonging to the Coronaviridae family, finally defined as “severe acute respiratory syndrome coronavirus 2” (SARS-CoV-2) for high sequence identity (i.e. up to 80%) with the homologous virus which caused the SARS outbreak in 2003 (i.e. SARS-CoV) [1]. At the time of writing this article, SARS-CoV-2 has already infected over 115,000 people in more than 115 different countries, causing nearly 4000 related deaths [2]. Structural analysis shows that SARS-CoV-2 probably derives

from a bat SARS-like coronavirus, which has been then transmitted to humans after emergence of mutations in the spike glycoprotein (protein S) and nucleocapsid N protein [3]. The mutation that occurred in the former protein is especially important, whereby viral spike glycoprotein mediates the entrance of the virus into the cell through cell receptor binding and membrane fusion. On the other hand, the N protein regulates the process of viral replication, thus influencing transcription and assembly.

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Altogether, mutations in these two proteins would then explain the unique characteristics of SARS-CoV-2 compared to the original SARS-CoV, i.e. enhanced infectious potency in humans, combined with relatively mitigated pathogenicity. In support of the former aspect, the effective reproductive number (R; i.e. the average number of secondary cases per infectious case) has been estimated at 2.6 (credibility interval, 2.1–5.1) for SARS-CoV-2 compared to 1.1 for SARS-CoV, whilst the doubling time of the epidemic has also been calculated as 3.6 days (comprised between 1.0 and 7.7 days) compared to approximately 16 days for SARS-CoV [4]. As concerns the mortality, the World Health Organization (WHO) provides daily estimates, which are obviously in progress due to the ongoing epidemics, the last of which attests that the death rate of COVID-19 is ~3.9% in China (3123/80,904) and ~2.4% abroad (686/28,673; e.g. 5.0% in Italy, 366/7375), compared to ~9.6% (774/8098) for SARS and 34.4% for Middle East respiratory syndrome (MERS; 866/2519). Between 8 and 15% (depending on the geographical setting and individual characteristics) of all SARS-CoV-2 positive cases can be classified as severe or necessitating intensive care unit (ICU) admission. Although the mortality rate of COVID-19 seems hence for now lower than that of SARS or MERS, the number of patients needing urgent critical care is remarkably larger than that of the two previous viral outbreaks, and may foster the collapse of local health care.

This is not the first case, nor it will probably be the last that a viral outbreak has become a public health concern, though COVID-19 displays distinctive features compared to previous coronavirus epidemics such as SARS and MERS, in that the pathogenicity of SARS-CoV-2 seems for now lower and the incubation, longer (usually up to 2 weeks), so the risk of contagion is magnified and the number of cases (and deaths) grows exponentially [5]. This is not really surprising if we look at the future mortality projection of the WHO between the years 2016 and 2060 [7], whereby the number of deaths for lower respiratory infections is expected to increase by over 50% during the next 40 years (i.e. from 2.96 to 4.62 million deaths per year).

This notable increase in mortality for pneumonia and other lower respiratory infections predictably encompasses also those caused by coronaviruses, as interstitial pneumonia – evolving toward acute respiratory distress syndrome (ARDS) in 10–15% of cases – is the most frequent and severe complication of SARS-CoV-2, which can then be followed by the onset of viral sepsis, disseminated intravascular coagulation (DIC) and multiorgan failure (MOF) [8, 9]. One

obvious question that arises here is whether, and eventually how, laboratory diagnostics could efficiently contribute to counteract this and other (future) viral outbreaks.

In Monitoring Etiological Response

The etiological diagnosis of SARS-CoV-2 is the first and most obvious setting where laboratory diagnostics plays an essential role. Both the WHO and the US Centers for Disease Control and Prevention (CDC), along with other national and international scientific organizations, have timely released detailed information for in-house development of reverse transcription-polymerase chain reaction (RT-PCR) tests, which have hence been straightforwardly implemented by many reference laboratories worldwide [11], and are now undergoing clearance by many regulatory agencies.

A crucial aspect that shall be underscored here is the need for developing rapid and effective communication among distant laboratories and research centers.

The enormous diffusion of this virus, with positive cases identified in over 115 worldwide countries and in almost every continent, highlights the vital need for developing diagnostic workflows even in the lack of physical sources of viral genomic nucleic acid, whereby the chance that outbreaks will spread to distant countries and become pandemic diseases is increasingly more likely, as recently demonstrated by COVID-19. Notably, the willingness of Chinese scientists to rapidly share genomic information has enabled to develop RT-PCR assays even before SARS-CoV-2 started to circulate in many countries, thus providing a timely and effective diagnostic response to a probable health crisis [12].

Nevertheless, as the ample volume of tests needed to face a large outbreak such as that sustained by SARS-CoV-2 overcomes the throughput capacities of single facilities, whilst shipment of samples toward reference laboratories is an important cause of diagnostic delays, the availability of commercial diagnostic kits in peripheral centers shall be part of the strategy for early and accurate identification of the largest possible number of infected patients.

Despite the obvious emergency to promptly develop efficient diagnostic tools, a thorough analytical and clinical validation of commercial RT-PCR tests before their introduction into the market and usage in clinical laboratories remains indispensable. Otherwise, the risk of generating false-negative (or positive) test results may undermine the huge efforts made by healthcare authorities for containing the outbreak.

In Monitoring Patient Response, Severity and Recovery

The second essential contribution that laboratory medicine could provide in the diagnostics of 2019-nCoV infection encompasses staging, prognostication and therapeutic monitoring of COVID-19. Not only RT-PCR tests will be vital for verifying the course of the infection, as well as the possible presence and extent of viremia, but many other laboratory tests may help assessing disease severity and predicting the risk of evolution toward ARDS, DIC and/ or MOF. A systematic literature review, which has recently been published [14], has highlighted the most important abnormalities observed in patients with COVID-19, 'mostly encompassing lymphopenia, increased values of C reactive protein (CRP), lactate dehydrogenase (LDH), erythrocyte sedimentation rate (ESR) and D-dimer, along with diminished concentration of serum albumin. Even more importantly, a number of hematological parameters were also found to predict progression toward severe or critical forms of COVID-19, including leukocytosis, neutrophilia and lymphopenia. In addition, an innovative parameter called MDW (monocyte volume distribution width - DxH 900 hematology analyzer, Beckman Coulter, Brea, CA, USA) was found to be significantly increased in all COVID-19 patients, especially those with worst clinical conditions (personal data, not shown). For prognostication purposes, also increased values of LDH, aspartate aminotransferase (AST), alanine aminotransferase (ALT), total bilirubin, creatinine, cardiac troponins, D-dimer, prothrombin time (PT), procalcitonin and CRP, together with decreased values of serum albumin, have been found of value.

The importance of hemostasis tests has then been emphasized in another study [15], including 94 patients with COVID-19, and showing that PT and D-dimer are significant predictors of disease severity. This finding not only supports the pivotal role of hemostasis testing in severe and/or systemic infectious diseases [16], but also confirms that consumption (disseminated) coagulopathy may be one of the most severe complications of patients with COVID-19.

These findings have also been confirmed in a subsequent study, which pooled data of 1099 patients with laboratory confirmed SARS-CoV-2 infection from 552 hospitals in 30 Chinese territories [17], demonstrating that COVID-19 patients have lymphopenia (83.2%), thrombocytopenia (36.2%), increased values of CRP (60.7%), LDH (41.0%), AST (22.2%), ALT (21.3%) and D-dimer (43.2%). In keeping with previous findings, the most predictive parameters of severe COVID-19 disease were lymphopenia (96.1% vs. 80.4%; odds ratio [OR], 5.96; 95% confidence interval [CI], 2.58-13.75), thrombocytopenia (57.7% vs. 31.6%; OR, 2.96;

95% CI, 2.07-4.22), leukocytosis (11.4% vs. 4.8%; OR, 2.54; 95% CI, 1.43-4.52), increased values of CRP (81.5% vs. 56.4%; OR, 3.40; 95% CI, 2.15-5.40), procalcitonin (13.7% vs. 3.7%; OR, 4.14; 95% CI, 2.06-8.33), LDH (58.1% vs. 37.2%; OR, 2.13; 95% CI, 1.45-3.14), AST (39.4% vs. 18.2%; OR, 2.92; 95% CI, 1.97-4.34), ALT (28.1% vs. 19.8%; OR, 1.59; 95% CI, 1.04-2.43) and D-dimer (59.6% vs. 43.2%; OR, 1.94; 95% CI, 1.27-2.97), whilst the median hemoglobin value was also found to be lower in patients with severe COVID-19 (128 vs. 135 g/L; $p < 0.01$). Each of these prognostic parameters retain a specific clinical and biological significance, which, altogether, can contribute to reflect the evolution toward more unfavorable clinical situations.

In Epidemiological Surveillance

Though not less, essential support given by diagnostic testing to counteracting viral outbreaks is the at the dawn of the third millennium. Laboratory automation, availability of high-throughput instrumentation [21], along with lower number of employees and reduced healthcare funding (especially for public facilities) have all contributed to considerably reduce the flexibility to develop emergent responses [22]. The availability and extended use of point-of-care testing (POCT) devices shall be regarded as an additional useful tool during outbreaks and other biological hazards, especially those sustained by viral infections [23, 24].

The lesson learnt from the recent outbreak in China is indeed paradigmatic, whereby the currently available healthcare resources were totally insufficient to manage the impressive number of patients seeking care in Wuhan. The situation has forced public authorities to rapidly build an entirely new 645,000-square-foot hospital, with approximately 1000 beds, ICUs, isolation wards and even a clinical laboratory inside.

Therefore, additional strategies shall be envisaged. The first and perhaps most important lesson that policymakers and hospital administrators shall learn from COVID-19 is that continuing to cut down human and economic resources will then generate huge organizational issues when the entire system of care, including laboratory diagnostics, will be challenged by an enormously amplified volume of tests to manage emergent situations [25]. Existing laboratories may be asked to enhance their usual throughput and contextually reduce their turnaround time, but this may not be sustainable always and anywhere. Urgent personnel recruitment for managing an enhanced volume of serological or molecular tests

will be needed, and shall be arranged as soon as possible considering that hands-on training is necessary for those who lack direct experience or skills in virological assays. Healthcare staff may also be temporarily moved from one laboratory to another (e.g. from a biochemistry to a virology laboratory), and this may have an impact on the efficiency of the former facility to maintain the usual throughout and turnaround time for routine and urgent non-virological tests.

Therefore, establishment of an efficient network of regional clinical laboratories, involving those which are not directly challenged by the outbreak and where samples can be conveyed, is a feasible solution, provided that a straightforward regulation for specimen transportation and biosafety is set and monitored. This, in turn, highlights an unavoidable need to place major efforts for allowing better and wider harmonization of laboratory results and information, encompassing both analytical and extra-analytical issues [26–28].

A clear and bulletproof safety policy shall also be instructed and communicated to the laboratory staff, encompassing all those measures that need to be established for preventing the health risks caused by the microorganism causing the outbreak. Two final aspects that shall be clearly acknowledged include the possibility that the healthcare staff may be directly infected by the virus [30], along with the safety measures established by local authorities, which may limit human movements and derange public transportation [31]. Both these aspects may contribute to substantially magnify the shortage of staff inside and outside the laboratory during the outbreak, and lead the way to a final consideration about the compelling need to develop national plans for emergency preparedness, which not only encompass all the previously discussed aspects, but also consider to invest more money in temporary stationary laboratory facilities, equipped with all the necessary instrumentation and with trained personnel, which could be rapidly transported to the site of emergency for supporting local testing needs.

Last but not least, it is essential that the laboratory personnel be instructed to communicate test results to the appropriate stakeholders (i.e. to the people who are officially in charge of dealing with the outbreak), thus avoiding to spread information that could generate unjustified panic, or inappropriate reassurance, among the general population [32].

Conclusion

The managed care of patients with SARS-CoV-2 infection entails early identification, rapid isolation, timely establishment of infection prevention and control (IPC) measures, together with symptomatic care for patients with mild disease and supportive treatment for those with severe COVID-19. Several lines of evidence attest that previous viral outbreaks may have been characterized by retarded identification and delayed healthcare response [33], and COVID-19 shall not be considered an exception to this rule [34]. Awareness and preparedness to face highly contagious viral outbreaks, such as that sustained by SARS-CoV-2, become imperative for preventing the health system from being strained and laboratory services from collapsing. Irrespective of its inherent definition [10], it is now virtually incontestable that laboratory medicine will increasingly provide an essential contribution to the diagnostic reasoning, managed care and therapeutic monitoring of the vast majority of human diseases [35], thus including infectious diseases [36, 37] and COVID-19, which has now been defined as global health emergency by the WHO.

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