COVID-19 diagnostic process in mainland China: The math beyond pneumonia

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In late December 2019, a cluster of pneumonia cases of unknown cause was first noted in human patients in the city of Wuhan, Hubei Province, China. The patients with this emerging disease presented with clinical signs of viral pneumonia similar to those observed in severe acute respiratory syndrome (SARS). Chinese scientists, including those in the Shanghai Public Health Clinical Center, the Chinese Center for Disease Control and Prevention, and the Wuhan Institute of Virology, immediately took actions to probe for the obligated emerging pathogen(s) through next-generation sequencing. A novel Betacoronavirus was soon identified, and its genome sequence was first made accessible to the world on January 12, 2020. On February 11, 2020, this novel coronavirus was named SARS-coronavirus 2 (SARS-CoV-2) by the International Committee on Taxonomy of Viruses and the associated disease was named coronavirus disease 2019 (COVID-19) by the World Health Organization.

COVID-19 is highly contagious. It has been reported that SARS-CoV-2 had an estimated basic case reproduction number ($R_0$) as high as 3.15 before the implementation of the emergency response on January 23, 2020, in mainland China. SARS-CoV-2 viral load in the upper respiratory system is highly dynamic along the progression of COVID-19, and it exhibits significant variation in different patients, with peaks on the order of $10^8$ viral genome copies/mL in throat swabs and $10^{11}$ copies/mL in sputum samples. SARS-CoV-2 also showed differential tissue tropism in different infected individuals. Viral load could be higher in fecal samples than in nasopharyngeal swab samples in some cases, likely because of oral transmission. The current standard operating procedure in clinical diagnosis uses pharyngeal swabbing to collect samples from the upper respiratory system. Limitations in preparing viral genetic materials for PCR tests include timing of sample collection and anatomic locations of sampling, which can be further complicated by practical variations in technical operations such as those in shipping, storage, and material/reagent preparation. The genetic sequence of SARS-CoV-2 enabled the rapid development of point-of-care real-time PCR diagnostic tests specific for SARS-CoV-2. PCR tests were used as a criterion standard measure to confirm active SARS-CoV-2 infection. There was, however, a false-negative rate close to 40% because of various issues, most of which are associated with sample preparation. This 40% false-negative rate is the rate of PCR testing in diagnosing SARS-CoV-2 infection, but not the false-negative rate of PCR test kits. In a recent external quality assessment program directed by the National Center for Clinical Laboratories in China, SARS-CoV-2 PCR test kits approved by China’s National Medical Products Administration were reported with an overall 95.4% positive test result when the input viral genomic load was 3200 copies/mL. The PCR test kit developed by the Beijing Genomics Institution detected 100% of the positive samples when the input was 3200 copies/mL and 97.1% of those with input level at 640 copies/mL; this kit was approved by the US Food and Drug Administration through emergency use authorization, and it became available for COVID-19 diagnosis in the United States on March 26, 2020.

There is a broad spectrum of COVID-19 symptoms, including fever, dry cough, shortness of breath, sputum production, diarrhea, fatigue, and others, most of which are associated with the respiratory and gastrointestinal systems. In mainland China, about 80% of the patients with confirmed infection have mild symptoms and 2.3% have fatal symptoms, whereas 1.2% are asymptomatic (calculated on the basis of data reported before February 11). The reported asymptomatic patients here were defined by a positive PCR test result accompanied by a lack of symptoms clinically, as described by patients, measured by doctors, and visualized by imaging, and the patients did not develop any symptoms after a fortnight quarantine. Patients who are still in the incubation period may remain asymptomatic in their upper respiratory system for weeks, although signs of pneumonia might have been developing in their lower respiratory system. There is evidence that infected individuals can be contagious even when they show no symptoms, and light or mild cases can develop into severe and even critical cases after a long period of time on account of lack of rest, mental stress, and/or malnutrition. An early and accurate diagnosis is extremely critical for containment and disease intervention, as well as for epidemiologic investigation.

Some patients in whom the virus was suspected had a clear history of contacts with confirmed and symptomatic patients but tested negative by PCR multiple times, whereas their symptoms varied from signs of viral pneumonia to mild or even no symptoms. Some patients had a documented history of exposure and showed typical signs of ground glass opacity in computed tomography (CT) scans; these patients were immediately admitted and treated as having COVID-19 with clinical diagnosis in China, with most of them testing positive by PCR and/or...
antibody assays during disease progression and recovery. Because of variations in the viral load of different individuals, and timing and anatomic locations of sampling for PCR, some infected patients were tested more than once but never had a sample with viral load that passed the PCR detection threshold. These patients with COVID-19 could have fully recovered without any positive PCR test results. Their immune systems, however, kept the footprints of the infection, virus-specific antibodies. By testing for SARS-CoV-2–specific antibodies, it was possible to determine whether the patients in whom the virus was suspected had a history of infection, even if the virus had been cleared after recovery. To obtain a more accurate number of COVID-19 cases, the Chinese Center for Disease Control and Prevention, along with scientists and doctors, developed a standard operating procedure for screening patients in whom the virus was suspected and confirming cases involving patients with infection who showed negative results in PCR tests by using SARS-CoV-2–specific antibody assays. A combination of information on exposure history, clinical presentation visualized on CT scans, presence of viral genes, and presence of virus-specific antibodies was used for diagnosing COVID-19 in mainland China. Of note, in an antibody (IgG) test trial conducted with >600 serum samples collected from >300 patients with “clinically diagnosed” COVID-19, more than 95% of the samples collected 21 to 25 days after disease onset showed strong positive results (An et al, data not shown).

The schematic flowchart delineating the standard operating procedure of the COVID-19 diagnostic process in mainland China is depicted in Fig 1. In brief, patients in whom COVID-19 is suspected and who have exhibited any COVID-19 symptoms and/or a documented history of contact with individuals with a confirmed case of COVID-19 were examined with CT scanning and PCR testing. Patients were then divided into 3 categories on the basis of findings in lung imaging. Signs of pneumonia in imaging and a positive PCR test result were used to diagnose and confirm COVID-19. Patients who had a contact history and typical signs of pneumonia in imaging but had a negative PCR test results were clinically diagnosed as having COVID-19 and subsequently retested with PCR and antibody assays. Before February 11, 2020, asymptomatic cases were reported as COVID-19 in mainland China. In early February, the 4th edition of Diagnosis and Treatment Protocol for Novel Coronavirus Pneumonia was released in mainland China; it recommended reporting SARS-CoV-2 infection in 2 categories: (1) COVID-19 (cases with confirmed infection and disease symptoms) and (2) asymptomatic infection. Any symptoms, including discomfort reported by patients, disease parameters measured by doctors (eg, high body temperature and low fingertip pulse oximeter reading), and signs of pneumonia in CT images, will define a case as symptomatic. The current reported COVID-19 cases in mainland China include all cases of COVID-19 (cases involving patients with infection plus symptoms) and asymptomatic cases (cases involving patients who never had symptoms) of SARS-CoV-2 infection determined before February 11, 2020.
COVID-19. In February, in Hankou Hospital in the city of Wuhan, an antibody test trial with serum samples collected from more than 300 “clinically diagnosed” patients showed that over 95% of these patients had a history of infection (An et al, data not shown); therefore, it is unlikely that clinical diagnosis would lead to an overestimation of COVID-19 cases in mainland China. On the basis of this scheme of diagnostic process as analyzed and published by the Chinese Center for Disease Control and Prevention, by February 11, 2020, there were 72,314 patients with COVID-19 reported in mainland China, of whom 1.2% were asymptomatic, 61.8% showed signs of pneumonia with a positive PCR result, and 14.6% were clinically diagnosed (Fig 1). In early February, mainland China started to report asymptomatic cases of SARS-CoV-2 infection and COVID-19 (infection plus disease symptoms) separately, according to the fourth edition of the Diagnosis and Treatment Protocol for Novel Coronavirus Pneumonia. Antibody assays were adopted to retest suspected cases involving patients who tested negative by PCR assay, and by March 31, 2020, a total of 81,554 confirmed cases of COVID-19 in mainland China were reported by the National Health Commission of China.

The goal of this Editorial is to provide a concise and clear flowchart to depict the COVID-19 diagnostic process developed and implemented in mainland China. We hope that this helps our readers better understand how the math was done to determine the number of reported cases of SARS-CoV-2 infection in mainland China. The math that results in the reported number of confirmed cases in mainland China is clearly beyond pneumonia. Despite our general understanding that COVID-19 is an acute respiratory disease caused by a novel coronavirus, clinical diagnosis was often complicated by clinical presentations, individual variations in patients and medical/laboratory operators, and the sensitivity and specificity of the tests. This diagnostic process used in mainland China may not be directly applicable to other areas in the world during the COVID-19 pandemic; however, it exploited current knowledge of epidemiology, disease characteristics, viral genetics, and immunology about COVID-19, and it provides an example of a comprehensive diagnostic approach.

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