Advances in Diagnostic Approaches of SARS-Cov-2 Virus

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Abstract

COVID-19 is caused by the SARS-CoV-2 virus, first identified as viral pneumonia, and later became a worldwide pandemic situation in December 2019. At present, there are 3,29,68,853 confirmed cases all over the world and 9,95,836 deaths (27th September 2020; 1:08 pm), USA is first on the list of confirmed and death cases. It can cause ARDS which is the worst-case scenario where the oxygen level is dangerously less and patients are kept under ventilation. ‘Prevention is better than cure’, there is no effective vaccine productions and specific therapeutics options are not standard in many countries. Personal hygiene and sanitization can help prevent this nCoV. Diagnostic methods can help detect the virus so that recommended measures can be taken to safeguard the patient’s life. Scientists are researching for future therapeutic options and vaccine production, also many clinical trials are going on for detecting the drug’s efficacy in viral load is reduced in SARS-CoV-2 patients.

Keywords

SARS-CoV-2, Prevention, Diagnostic approaches

Abbreviations

COVID-19: Corona Virus Disease 2019; ARDS: Acute Respiratory Distress Syndrome; QRT-PCR: Quantitative Real Time-Polymerase Chain Reaction; CT: Computed Tomography; LUS: Lung Ultrasound; AI: Artificial intelligence

Introduction

The COVID-19 was first likely to have originated in the Huanan seafood market in Wuhan city Hubei province, Central China, even though the initial source of origination is not known still. It was first recorded as viral pneumonia and later it became a world-wide pandemic situation. Corona viruses fall under the Corona viridae family and Ortho Corona viridae subfamily [1]. It causes infections like a common cold, respiratory problems, and more severe problems like SARS and MERS and its main source of spread is the respiratory tract. Elderly people are likely to be more associated with ARDS and cytokine storms [2]. This virus has the potential to grow and become a world-wide pandemic, on 28th of February 2020 a very high-level threat to the CoV epidemic has been raised by the WHO. SARS-CoV-2 transmission is faster and higher compared to previous viruses [3]. Through small droplets from the nose or mouth, it can easily spread from person-person through sneezing, coughing and speaking. Among 1 out of every 5 people is affected by a corona virus and becomes severely sick [4].

Prevalence

The outburst of COVID-19 pandemic originated from Huanan seafood market, China. As per WHO, the samples taken from the surroundings of this market were marked positive for this infection. Yet the interrelations among the infections with distinct animals have not been identified but the previous record says that bats are the host for more than 30 infections which concludes that it might be a source for COVID-19 [5,6]. On 11th of March, 2020 considering the uplift in severity and unfavorable consequences of the infection, WHO proclaimed it as an overall pandemic [7]. It has been retrieved from the WHO; up to 27th of September 2020 the complete count of positive cases registered worldwide was 3,29,68,853 and the worldwide death rate rose up to 9,95,836. As per the evidence provided by WHO, as of 27 September 2020 USA stands first in position worldwide and contributes to about 21.25 % of total positive cases followed by India (18.42%), Brazil (14.31%), Russia (3.49%) & Columbia (2.44%). Countries like Maldives (0.03%), Myanmar (0.028%) Zimbabwe (0.024%), Thailand (0.0108%) are least affected and stand down in position is illustrated in the below (Figure 1 and Figure 2).

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Mechanism

The ingress of SARS-COV-2 into the cell which is being encountered commences with the synergy of Spike proteins (S-Proteins) made up of Glycoprotein with the receptors of the host. As per the types of viruses, the sites of Receptor Binding Domains (RBD) within the S1 locale of a COVID S protein differ. Some may have RBD at N-terminus for e.g. MHV and some may have such at C-terminus as in the case of SARS-COV [8,9]. At that point, viral particles access over host cell cytosol which can generally be achieved by Cathepsin or other proteases like TMPRSS2 etc. This helps in cleaving S-protein, which usually gets cleaved at 2 different sites of which the first cleavage is vital for division of Receptor Binding Domain & fusion domains of S protein. Subsequently, the second cleavage for revealing the fusion peptides [10,11], after the cell entry, the RNA gets uncoated and the replicase gene on the RNA strand is translated into two replicase poly proteins, further the proteinases of virus process these poly proteins to obtain independent proteins of replicate enzymes. These proteins bring about the output of full-length negative sense RNA and it functions as positive sense RNA template in future and the full-length negative sense RNA is reproduced to deliver shorter mRNAs [12].

Diagnostic Approaches - Molecular Aspect

Quantitative real-time reverse transcription-Polymerase chain reaction (QRT-PCR)

One of the most common and promising methods of diagnosis used in the detection COVID-19 is Quantitative Real-Time Reverse Transcriptase-Polymerase Chain Reaction [13]. The steps involved in this diagnostic method are as follows: 1) Extraction of RNA samples from the patients; 2) The RNA extracted is converted into cDNA by reverse transcription and it is amplified by specific fragments of cDNA by PCR; 3) Fluorescent signals are studied. When the Cycle threshold (C) values are lesser, it is an indication of diagnostic results as positive. Generally, it takes about 2-4 hour’s when performed under trained people in a clinical laboratory whereas; automated QRT-PCR is expected to give the result in less than an hour. This method is very useful for diagnosis in early stages compared to antigen-antibody reaction [14]. If there is any

![Figure 1: Number of positive cases according to each country till 27th September 2020.](image1)

![Figure 2: Number of death cases according to each country till 27th September 2020.](image2)
other material that is involved it will affect the results and will show as “false-positive results. Moreover, micro fluidic digital array PCR helps to lower the false-positive rate because it gives gene qualification at single molecule level [15].

Detection by CRISPR/CAS Technology

CRISPR/Cas system is a kind of impressive genetic engineering tool that mainly focuses on editing the desired genome of interest [16,17]. These technologies are now freshly introduced in detecting the viral particles with more promising results [18,19]. Major components of this technology include the CAS proteins which give the enzymatic apparatus needed to the procurement of new spacers from, targeting, attacking components, the guide RNAs helps in target sequence selectivity for the system and the fragments of target DNA [20,21]. The Cas proteins like 12a & 13a results in the guaranteed cleavage when bound over the target, in which a nonspecific single stranded DNA or RNA marked with fluorescent dye gets digested and released out [22,23]. Recent studies say that these CRISPR/Cas 13a tests alongside with isothermal amplification can locate SARS-COV-2 which was affirmed already by the US FDA for crisis use [24,25]. Also, the combination of cas12a with RT- LAMP shows a greater relevance and sensitivity over the samples [26]. Therefore, this technology is assessed as a promising diagnostic method for the detection of SARS-COV-2, since it can be incorporated with any nucleic acid augmentation frameworks [27,28].

Detection by genome sequencing

Genome sequencing is a kind of methodology in which the arrangement of DNA nucleotides or the bases in the whole genome and the composition of double helix DNA are unraveled [29]. It is notable for its agility and precision in recognizing the unknown viral particles entering our body and also it has a numerous point of interest in virus detection. To start with, it empowers examination of the SARS-CoV-2 development during transmission and following of COVID developments [30]. It is then followed by viable prohibition of other microbes. Also, genomic sequencing can distinguish numerous microorganisms existing in a solitary patient, which helps execute sensible medications but requires lavish equipment and skilled personnel [31]. Higher recognition efficiency can be achieved by using droplet based single cell sequencing technology. At present, this method has first concern at the ongoing phase of the epidemic, yet patient’s genome sequencing will provide a better understanding about the human susceptibility over the SARS-COV-2 infection [32]. By interpreting the genomic sequences of the affected patients, the transmission dynamics of SARS-COV-2 can be very well understood which aids in the rapid development of vaccines against this infection and their diagnostic approaches is illustrated in (Figure 3).

Imaging Examination

Chest computed tomography (CT)

It is an x-ray imaging technique where narrow x-ray beams are focused on patients’ body; it gives the cross-sectional images of the required target [33]. In this aspect, it is focused on cross-sectional images of the lungs by CT where we get more details about radiographic features in most of the COVID-19 patients [34]. CT scans are strongly recommended to use as a diagnostic option, there are five stages classified based on CT imaging stages [35,36] is presented in (Table 1).

Lung ultrasound (LUS)

Ultrasound is used to image the internals of the body; it is done by the production of sound waves by probes or transducers above human hearing threshold frequencies, these probes can also be sent inside the body to get more detailed image [37]. LUS is an easy option for diagnosis of suspected COVID-19 patients because, it can be done at any stage of diagnostics or therapeutic processes while the patients still
on bedside [38]. There are some advantages compared to CT technique, LUS has no radiation, it is low cost and can be performed repeatedly faster [39,40]. LUS has limitations when compared to CT; low sensitivity, lesions that are presented deep and intrapulmonary cannot be detected [41].

**Treatment**

There is no particular therapeutic option available for COVID-19 and vaccines are not yet discovered, scientists are undergoing research for developing vaccines and having treatment plans for future aspects [42]. vaccine as an early preventative strategy for SARS-CoV-2 virus. There are many antiviral drugs like hydroxychloroquine and Lopinavir/Ritonavir were taken by people for treatment [43]. There are some selective treatment options based on the 7th edition of ‘COVID-19 diagnosis and treatment guideline’ released by NCH of the people’s Republic of China, it is based on three categories are as follows: 1. Standard treatment; 2. Treatment for severe and critical cases; 3. Traditional Chinese medicine [44].

**Conclusion**

SARS-CoV-2 is a highly infectious and transferrable virus which causes severe conditions, even though there are no therapeutic options and vaccines available. Diagnostics methods (Molecular aspect and Imaging examination) are there to detect the virus but, improvements and advancements are much needed to faster the rate of diagnosis. The use of AI technology can also be helpful in diagnostic aspects like; robots can be programmed for collecting samples which reduces human force. In near future, the advanced treatment options of vaccines production and molecular diagnostic methods will provide a great help for disease diagnosis and detection.

**Acknowledgement**

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**References**


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**Table 1: Stages based on CT image in SARS-CoV-2 virus.**

<table>
<thead>
<tr>
<th>S.No</th>
<th>Stages based on CT image</th>
<th>Conditions according to each stage</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>Ultra-early stage</td>
<td>No clinical manifestation in this stage, laboratory results are negative but tests positive for throat swab generally between 1-2 weeks after exposure to contaminated areas.</td>
</tr>
<tr>
<td>2</td>
<td>Early stage</td>
<td>Initial period within three days once encountered with cold, wheeze, dry cough, pyrexia etc., it shows particular numerous strewed irregulars or groups of ground-glass opacities, and divided by beehive-like stiffen interlobular septa.</td>
</tr>
<tr>
<td>3</td>
<td>Rapid Progression stage</td>
<td>Period of three to seven days after being influenced by the virus, a merged &amp; huge-scale light coalesce with air-bronchogram inside was manifested by CT.</td>
</tr>
<tr>
<td>4</td>
<td>Consolidation stage</td>
<td>Period of about one to two weeks after exposure to clinical manifestation, CT image portrayed several strewed coalesce in lower density and lesser range than the before stage.</td>
</tr>
<tr>
<td>5</td>
<td>Dissipation stage</td>
<td>Hardly between 2-3 weeks after clinical manifestation, furthermore was the range of lesions reduced. Patchy consolidation or strip-like opacity was shown by CT image, after sometime grid-like denser interlobular septum, thickening and strip-like twist of bronchial wall and less scattered patchy consolidations was observed.</td>
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</tbody>
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