A laboratory-based study of COVID-19 in Casablanca, Morocco

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Abstract

Background: Given the spread of coronavirus disease 2019 (COVID-19) and its impact on human health, laboratory confirmation of diagnosis is essential.

Objective: This study examined the contribution of laboratory diagnosis to the detection of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) in the diagnosis of COVID-19, taking into account patient risk of exposure to SARS-CoV-2, clinical symptoms and comorbidities.

Methods: A cross-sectional, laboratory-based study was carried out from 1 April 2020 to 30 April 2020 at the National Reference Laboratory in Morocco using nasopharyngeal samples from patients admitted to the Cheikh Khalifa International University Hospital or other hospitals in Casablanca. A one-step reverse transcription real-time polymerase chain reaction (RT-PCR) was used to detect the presence of the SARS-CoV-2 genome. A national epidemiological investigation form was used to analyze patient exposure risk, clinical symptoms and comorbidities.

Results: A total of 793 samples from 375 patients were analyzed and 1150 RT-PCR tests were conducted; 116 patients (30.93%) were COVID-19 positive. Travel to a risk zone, contact with a confirmed COVID-19 case and contact with a person who had been in a risk zone were significantly associated with being positive for COVID-19. Fever and cough were the main symptoms; 7.76% of positive patients were asymptomatic.

Conclusion: This is the first laboratory-based study in Morocco for the diagnosis of COVID-19. Laboratory diagnosis of COVID-19 by RT-PCR associated with knowledge of exposure risk factors and clinical symptoms and comorbidities remains essential for clinicians for early, appropriate medical management COVID-19 patients.
Introduction

In mid-December 2019, an epidemic of a viral respiratory disease, officially named by the World Health Organization (WHO) as coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), was discovered in the city of Wuhan, Hubei province, China. This epidemic then spread to the rest of the world, prompting WHO to declare a public health emergency of international concern and subsequently to declare the COVID-19 outbreak a pandemic (1, 2).

Since the start of COVID-19 in December 2019 to April 30, 2020, a total 3 090 445 confirmed cases of COVID-19 have been reported worldwide, with 217 769 deaths. As of April 30, 2020, there were 182 417 confirmed cases in the Eastern Mediterranean region with 7447 deaths. In the WHO African region, a total of 24 713 confirmed case with 938 deaths were reported. In Morocco in April 30, 2020, there were 4359 confirmed cases with 168 deaths was reported (1, 3). At present, there are no specific antiviral drugs or vaccine against COVID-19 infection for potential therapy of humans. Studies have shown that broad-spectrum antivirals like remdesivir, oseltamivir, lopinavir, ritonavir, ganciclovir and the antiparasitic chloroquine could be highly effective in the control of COVID-19 (4, 5).

Coronaviruses are a large family of viruses that can be pathogenic in humans and animals. The viruses of the Coronaviridae family are enveloped viruses with a single-stranded RNA genome of positive polarity (6). Recently, Three types of coronaviruses, which are not found naturally in humans but in mammals, have been discovered and have been the origin of serious lung infections (viral pneumonia): SARS-CoV, Middle east respiratory syndrome coronavirus (MERS-CoV) and SARS-CoV-2, responsible respectively for severe acute respiratory syndrome (SARS), Middle East respiratory syndrome (MERS) and COVID-19 (4, 7-9). For the last coronavirus responsible for COVID-19, the most common symptoms are fever, fatigue and a dry cough. Some patients experience pain, nasal congestion, runny nose, sore throat or diarrhea. About one in six people with the disease have more severe symptoms, including dyspnea. Seniors and those with other health conditions (high blood pressure, heart problems or diabetes) are more likely to have serious symptoms (9, 10).

Currently, the diagnosis of SARS-CoV-2 infection is a public health issue. SARS-CoV-2 screening allows the diagnosis of COVID-19 infection. This screening is based on the detection of the RNA of SARS-CoV-2 by a reverse transcriptase PCR technique in real time. These PCR assays specifically detect specific genes of SARS-CoV-2. The most relevant targets are: the envelope (E gene) gene, the RNA dependent RNA polymerase (RdRp gene) and the nucleocapsid (N gene) and the ORF1ab encoding proteins for RNA replication (ORF1ab gene) (11). Alongside the specific RT-PCR, the diagnosis of SARS-CoV-2 infection may also be made by the determination of IgM / IgG in serum (12). In Morocco several laboratories have been authorized by the Ministry of Health to carry out these diagnostic tests. Among these laboratories, the National Reference Laboratory (Laboratoire national de reference, LNR) attached to the Mohammed VI University of Health Sciences is an advanced laboratory equipped with several molecular diagnostic platforms, including those for COVID-19.

In Morocco, there no data on laboratory-based studies of the management of COVID-19. Given the spread of the new coronavirus and its impact on human health, laboratory confirmation of diagnosis is essential. This study aims to provide the contribution of RT-PCR to the detection of SARS-CoV-2 in the diagnosis of COVID-19 from patients taking into account the patient exposition risk to SARS-CoV-2, clinical symptoms and comorbidities.

Materials and methods

Ethical considerations

This study was approved by the Ethical Committee for Biomedical Research of Mohammed VI University of Health Sciences, Casablanca, Morocco (Ref: CE_UM6SS/2/06/2020 – May 4,
The patients or legal guardians were informed in writing about the study, and approved and consented to the use of their data for scientific purposes. The data were systematically entered in the LNR system database guaranteed by a restricted access code, and the study was carried out confidentially and anonymously.

**Patients, sampling and study design**

From 1 April 2020 to 30 April 2020, we carried out a cross-sectional, laboratory-based study that included samples from patients admitted to the Cheikh Khalifa International University Hospital or from to other hospitals in Casablanca, Morocco. All samples were collected for COVID-19 diagnosis. Sampling was done using nasopharyngeal swab conserved in universal transport media (UTM) for viruses. The collected swabs were placed into 3 ml of UTM and sent to the LNR in a triple packaging system at + 4 °C. All laboratory procedures were conducted at the LNR.

To study the exposure risk factors to SARS-CoV-2, clinical symptoms and comorbidities, we used the national epidemiological investigation form for possible cases of COVID-19 provided by the Moroccan Ministry of Health. All samples were accompanied by this completed form. We investigated data related to: (1) identity (gender, age, sex and address), (2) risk factors for exposure to COVID-19 (travel to a risk zone, contact with a confirmed case, attendance at a hospital structure, contact with a person who had visited a risk zone, and is a patient of a healthcare professional working in the hospital or in a laboratory realizing COVID-19 diagnostic test, (3) clinical symptoms (start date of symptoms, fever, cough, sore throat, breathing difficulty, headache, myalgia, diarrhea, abdominal pain, vomiting, respiratory distress and other non-specific symptoms), (4) co-morbidities and risk factors associated with COVID-19 (diabetes, high blood pressure, chronic respiratory disease, pregnancy, immunosuppression and other non-specific comorbidity). All samples sent to LNR without the completed form were excluded from the study.

**RNA extraction and RT-PCR SARS-CoV2 detection**

For RNA extraction, we used AccuPrep® Viral RNA Extraction Kit (Bioneer Corporation, Munpyeongseo-ro, Daedeok-gu, Daejeon, Republic of Korea) according to the manufacturer's instructions. A one-step reverse transcription real-time polymerase chain reaction (RT-PCR) was used to detect the presence of the SARS-CoV-2 genome by amplifying specific RdRp, E, and N genes using GeneFinder™ COVID-19 Plus RealAmp Kit according to the manufacturer's protocol (OSANG Healthcare Co., Ltd, Anyangcheondong-ro, Dongan-gu, Anyang-si, Gyeonggi-do, Korea). All RT-PCR assays were performed on a CFX96 Real-Time System (Bio-Rad, Hercules, California, United States). For test interpretation, we used two recommendations: (1) the recommendations of GeneFinder™ COVID-19 Plus RealAmp Kit according to the manufacturer's protocol, (2) recommendations for the interpretation of laboratory tests and recommendations for laboratories investigating SARS published by WHO (13) (Table 1).

**The laboratory workflow for the confirmation of COVID-19 cases**

For confirmation of COVID-19 cases, our laboratory used validated commercial kit that detect three targets of SARS-CoV-2 genome. Of these three target, N and RdRp genes are the most specific to SARS-CoV-2. Therefore, the presence of one and/or the two genes are sufficient the confirmed a COVID-19 cases. To avoid samples and PCR contamination, with the Biological Safety Levels 3 environment for RNA extraction and handling primary samples from suspect patients, all the further step were handled separately: oligonucleotides and PCR mastermix reagents were handled in one room under a PCR hood with specified laboratory coats. Addition of sample RNA to the mastermix took place in a second room. The third room is the machine
room to launch the PCR on CFX96 Thermal cycler. This workflow takes into account of international standards for the organization of laboratories practicing the diagnosis of covid-19 (11, 13)

Statistical analysis
Anonymous computerized data were analyzed by EpiInfo 7 (Centers for Disease Control, Atlanta, Georgia, United States) and Microsoft Excel (Microsoft, Morocco). Qualitative variables were analyzed by performing non-parametric tests: The chi square test or Fisher's exact test or Yates correction test. The estimation of associations between COVID-19 and exposure risk factors, clinical symptoms and comorbidities were assessed by calculating odd ratios (OR) or risk ratios (RR). Differences were considered significant if the p-value was < 0.05. Statistical significance was determined with 95% confidence internals (CI).

Results
During the study period, 793 samples were analyzed by RT-PCR from 375 patients. A total of 1150 RT-PCR tests were completed with a range of 1 to 3 tests per patient. Among the 375 patients, RT-PCR for the detection of SARS-CoV-2 was positive in 116 patients (30.93%). The sex ratio (Males/Females) of positive patients was 1.37 with an overall average age of 47± 20 years old (aged from 5 to 84 years).

Of all the exposure risk factors studied, travel to a risk zone, contact with a confirmed COVID-19 case and the contact with a person who has sojourned in a risk zone were significantly associated to the COVID-19 (Table 2). The main factor strongly associated with contracting COVID-19 was contact with a confirmed COVID-19 case. The risk factors relative to a risk zone were a country where an epidemic was declared or a national region where a high concentration of cases was detected.

As for the clinical symptoms of COVID-19 positive patients, the average time from symptom onset to the confirmation of COVID-19 by RT-PCR at the laboratory was 14.13±6.61 days (range: 3 to 30 days).

The main statistically significant symptoms of COVID-19-positive patients were fever and cough (Table 3). For patients with fever, the average temperature was 38.3±0.39°C. A total of 7.76 % of positive patients (9 patients) were asymptomatic. Moreover, several other nonspecific symptoms were associated with COVID-19-positive patients. Mainly, 8.62% of patients had anosmia, 4.31% had ageusia and 2.58 % had dizziness. Other nonspecific symptoms described among less than one percent of patients were anorexia, dyspnea, respiratory distress, nausea, vomiting and anemia.

In our study, age over 60 years was a statistically significant comorbidity factor (accounting for 38.83% of patients; RR = 1.44, 95% CI = 1.055 – 1.969, p-value = 0.02) associated with COVID-19 (Table 4). Others comorbidities that could potentially be associated with COVID-19 were presence of diabetes (12.05% of positive patients) and high blood pressure (19.28% of positive patients).

Discussion
Since the start of outbreak in December 2019, the SARS-CoV-2 virus has continued to spread worldwide (3). The widespread distribution of this virus, more than 213 countries and territories, has led to a major concern, globally (14, 15). After MERS-CoV and SARS-CoV, the SARS-CoV-2 virus is the third contagious coronavirus leading to an epidemic in the 21st century (9). The objective of the current study was to provide the contribution of PCR to the detection of SARS-CoV-2 in the diagnosis of COVID-19 from patients taking into account of the patient exposition risk to SARS-CoV-2, clinical symptoms and comorbidities. Laboratory diagnosis is necessary because it mainly distinguished COVID-19 from other known viruses of
pneumonia and from others non-infectious diseases. Identification of COVID-19 includes virus isolation and/or viral nucleic acid detection. In the present study, for only one month of April, we realized more than 1150 RT-PCR tests. The overall prevalence of COVID-19 was 30.93% taking into account our inclusion criteria. The investigation of exposure risk factors to COVID-19 of our study revealed that: travel to a risk zone, contact with a confirmed case and contact with a person who has been in a risk zone were the major risk factors exposure to COVID-19. This is also well described in the WHO suspect, probable and confirmed COVID-19 cases, as well as contact cases definitions, the risk factors found in our study constitute some of the main factors that contribute to the contamination of SARS-CoV2 and to developing COVID-19 (3).

The average time from symptoms onset to the confirmation of COVID-19 by RT-PCR at the laboratory was 14.13±6.61 days and ranged from 3 to 30 days. The mean incubation periods of COVID-19 in recent publications have differed slightly. Lauer et al., with 181 cases with confirmed SARS-CoV-2 infection detected outside Hubei province in China before 24 February 2020, reported that the median incubation period for these cases was 5.5 days (CI, 4.4 to 7.0 days) (16). Wang et al., in Wuhan, China with 138 cases, reported that the median durations from first symptoms range to were 5 days to 8 days (17). In summary, The incubation means ranged from 3 days to 14 days with an average CI of 0 to 30 days (10, 16, 17).

Importantly, 7.76 % of positive patients (9 patients) were asymptomatic. Elsewhere, the Epidemiology Working Group for NCIP Epidemic Response of the Chinese Center for Disease Control and Prevention, with a total of 72,314 patients, reported that there were 889 asymptomatic cases (1.2%), and 44,672 (61.8%) confirmed cases among the total number of patients (18). They had no symptoms upon confirmation of diagnosis of COVID-19 by RT-PCR. As reported elsewhere (19, 20), these asymptomatic patients constitute a big challenge for the management of COVID-19, because asymptomatic patients play a key role in the spread of virus.

The main significant symptoms associated with COVID-19 in our study were fever and cough. For patients with fever, the average temperature was 38.3±0.39. Patient age over 60 years was a significant comorbidity factor associated with the COVID-19 followed by diabetes and patients with high blood pressure. Out of these findings, several study reported common clinical symptoms include fever, cough, myalgia upper respiratory tract symptoms like rhinorrhea, sneezing, and sore throat and in addition to the increasing dyspnea with hypoxemia. Some patients infected with COVID-19 developed gastrointestinal symptoms like diarrhea (4, 21). Around the world, the most common symptoms were fever (98%), cough (76%), myalgia or fatigue (44%); and atypical symptoms included sputum (28%), headache (8%), hemoptysis (5%) and diarrhea (3%) (10, 21). As already reported, many underlying diseases (comorbidities) increase the risk of patients with COVID-19, including particularly elderly patient, diabetes, hypertension, but also cardiovascular disease (21).

Limitations
Because this study focused on laboratory-reported cases of COVID-19 and RT-PCR analysis, the evolution of clinical and biological parameters of patients was not assessed.

Conclusion
This is the first laboratory-based study in Morocco for the diagnosis of COVID-19. Through this study, we concluded that the contribution of laboratory for the diagnosis of COVID-19 cases by RT-PCR is essential for management of patients with COVID-19. We also argued that laboratory tests associated to the knowledge of exposure risk factors, clinical symptoms and comorbidities remains determining for clinicians in the early appropriate medical management of patients with COVID-19.
References

Table 1: Results interpretation of GeneFinder™ COVID-19 Plus RealAmp Kit according to
the manufacturer recommendations (March-2020 “rev.2” IFMR-45)

<table>
<thead>
<tr>
<th>Results Of RT-PCR</th>
<th>Ct&lt;sup&gt;a&lt;/sup&gt; range</th>
<th>RdRp&lt;sup&gt;b&lt;/sup&gt;</th>
<th>E&lt;sup&gt;c&lt;/sup&gt;</th>
<th>N&lt;sup&gt;d&lt;/sup&gt;</th>
<th>IC&lt;sup&gt;e&lt;/sup&gt;</th>
</tr>
</thead>
<tbody>
<tr>
<td>COVID-19 Positive</td>
<td>≤ 43  ≤ 43 ≤ 43 ≤ 35</td>
<td>≤ 43 ≤ 43 U.D ≤ 35</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>COVID-19 Positive if RdRp ≤ 43 after the second test</td>
<td>≤ 43 U.D U.D ≤ 35</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>COVID-19 Positive if E and N ≤ 43 after the second test</td>
<td>U.D ≤ 43 ≤ 43 ≤ 35</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>COVID-19 Positive if N ≤ 43 after the second test</td>
<td>U.D U.D ≤ 35</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Beta coronavirus</td>
<td>U.D ≤ 43 U.D ≤ 35</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Negative</td>
<td>U.D U.D U.D ≤ 35</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

<sup>a</sup> Ct: Cycle Threshold <br>
<sup>b</sup> RNA dependent RNA polymerase (RdRp gene) gene <br>
<sup>c</sup> Envelope (E gene) gene <br>
<sup>d</sup> Nucleocapsid (N gene) gene <br>
<sup>e</sup> IC: internal control <br>
<sup>f</sup> U.D = undetermined
Table 2: Exposure risk factors to COVID-19 of positive patients admitted to the Cheikh Khalifa International University Hospital in Casablanca, Morocco, from 1st to April 30th 2020.

<table>
<thead>
<tr>
<th>Risk factors exposure to COVID-19</th>
<th>Percentage of COVID-19 positive patients</th>
<th>Risk Ratio</th>
<th>CI 95%&lt;sup&gt;a&lt;/sup&gt;</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Travel to a risk zone</td>
<td>14.29</td>
<td>2.29</td>
<td>1.347 - 6.395</td>
<td>0.004</td>
</tr>
<tr>
<td>Contact with a confirmed COVID-19 case</td>
<td>66.67</td>
<td>1.55</td>
<td>1.255 - 1.922</td>
<td>0.0002</td>
</tr>
<tr>
<td>Attendance at a hospital structure</td>
<td>14.29</td>
<td>0.5</td>
<td>0.287 - 0.886</td>
<td>0.01</td>
</tr>
<tr>
<td>Contact with a person who has sojourned in a risk zone</td>
<td>29.76</td>
<td>1.81</td>
<td>1.169 - 2.826</td>
<td>0.008</td>
</tr>
<tr>
<td>Is the patient a healthcare professional working in the Hospital or in a laboratory realizing COVID-19 diagnostic test?</td>
<td>5.95</td>
<td>0.43</td>
<td>0.174 - 1.079</td>
<td>0.05</td>
</tr>
</tbody>
</table>

<sup>a</sup> CI: confidence interval
Table 3: Clinical symptoms of COVID-19 positive patients admitted to the Cheikh Khalifa International University Hospital in Casablanca, Morocco, from 1st to April 30th 2020.

<table>
<thead>
<tr>
<th>Clinical symptoms</th>
<th>Percentage of COVID-19 positive patients</th>
<th>Odd Ratio</th>
<th>CI 95%&lt;sup&gt;a&lt;/sup&gt;</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Fever</td>
<td>47.44</td>
<td>2.66</td>
<td>1.553 - 4.554</td>
<td>0.0002</td>
</tr>
<tr>
<td>Cough</td>
<td>53.86</td>
<td>2.36</td>
<td>1.397 - 4.003</td>
<td>0.001</td>
</tr>
<tr>
<td>Sore throat</td>
<td>20.51</td>
<td>0.93</td>
<td>0.4925 - 1.757</td>
<td>0.823</td>
</tr>
<tr>
<td>Breathing difficulty</td>
<td>23.08</td>
<td>0.66</td>
<td>0.363 - 1.203</td>
<td>0.173</td>
</tr>
<tr>
<td>Headache</td>
<td>37.18</td>
<td>1.48</td>
<td>0.861 - 2.558</td>
<td>0.153</td>
</tr>
<tr>
<td>Myalgia</td>
<td>30.77</td>
<td>1.37</td>
<td>0.777 - 2.431</td>
<td>0.273</td>
</tr>
<tr>
<td>Diarrhea</td>
<td>19.23</td>
<td>1.45</td>
<td>0.739 - 2.878</td>
<td>0.273</td>
</tr>
<tr>
<td>Abdominal pain</td>
<td>12.82</td>
<td>1.01</td>
<td>0.468 - 2.196</td>
<td>0.972</td>
</tr>
<tr>
<td>Vomiting</td>
<td>11.54</td>
<td>1.24</td>
<td>0.543 - 2.842</td>
<td>0.61</td>
</tr>
</tbody>
</table>

<sup>a</sup> CI: confidence interval
Table 4: Comorbidities factors of COVID-19 positive patients admitted to the Cheikh Khalifa International University Hospital in Casablanca, Morocco, from 1st to April 30th 2020.

<table>
<thead>
<tr>
<th>Comorbidities factors</th>
<th>Percentage of COVID-19 patients</th>
<th>Risk Ratio</th>
<th>CI 95%&lt;sup&gt;a&lt;/sup&gt;</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Diabetes</td>
<td>12,05</td>
<td>0,78</td>
<td>0,405 - 1,513</td>
<td>0,46</td>
</tr>
<tr>
<td>High blood pressure</td>
<td>19,28</td>
<td>1,15</td>
<td>0,678 - 1,955</td>
<td>0,6</td>
</tr>
<tr>
<td>Chronic respiratory disease</td>
<td>4,82</td>
<td>0,39</td>
<td>0,142 - 1,093</td>
<td>0,05</td>
</tr>
<tr>
<td></td>
<td></td>
<td></td>
<td>0,122 -</td>
<td></td>
</tr>
<tr>
<td>Pregnancy</td>
<td>1,2</td>
<td>1,33</td>
<td>14,488</td>
<td>0,81</td>
</tr>
<tr>
<td>Immunosuppression</td>
<td>1,2</td>
<td>0,89</td>
<td>0,094 - 8,413</td>
<td>0,92</td>
</tr>
<tr>
<td>Age over 60 years old</td>
<td>38,83</td>
<td>1,44</td>
<td>1,055 - 1,969</td>
<td>0,02</td>
</tr>
</tbody>
</table>

<sup>a</sup> CI: confidence interval