Brief communication

Definitive diagnosis in suspected Middle East Respiratory Syndrome Coronavirus cases

Elisa Rubio\textsuperscript{1,2}, Miguel J. Martínez\textsuperscript{1,2}, Verónica Gonzalo\textsuperscript{1,2}, Josep Barrachina\textsuperscript{1,2}, Núria Torner\textsuperscript{3}, Ana I. Martínez\textsuperscript{3}, Mireia Jane\textsuperscript{3}, Anna Vilella\textsuperscript{2,4}, Ana del Rio\textsuperscript{5}, Natalia Rodríguez-Valero\textsuperscript{2,6}, María Jesús Pinazo\textsuperscript{2,6}, José Muñoz\textsuperscript{2,6}, Alex Soriano\textsuperscript{5}, Antoni Trilla\textsuperscript{2,4}, Jordi Vila\textsuperscript{1,2}, and Ma Àngeles Marcos\textsuperscript{1,2*}

\textsuperscript{1}Department of Clinical Mirobiology, Hospital Clinic, University of Barcelona, Barcelona, Spain, \textsuperscript{2}ISGlobal Barcelona Institute for Global Health, Barcelona, Spain, \textsuperscript{3}Agencia de Salud Pública de Cataluña, Ciber de Epidemiología y Salud Pública CIBERESP, Spain, \textsuperscript{4}Public Health Department, Hospital Clinic, University of Barcelona, Barcelona, Spain, \textsuperscript{5}Department of Infectious Diseases, Hospital Clinic, University of Barcelona, IDIBAPS, Barcelona, Spain and \textsuperscript{6}Centre for Research in International Health (CRESIB), Hospital Clinic, University of Barcelona, Barcelona, Spain

*To whom correspondence should be addressed. Email: mmarcos@clinic.cat

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Abstract

We evaluated the microbiological diagnosis in 14 patients with epidemiological and clinical suspicion of Middle East Respiratory Syndrome Coronavirus (MERS-CoV) attended in a non-endemic area between June 2015 and January 2017. While no MERS-CoV was detected, other respiratory viruses were identified in 12 cases and Mycoplasma pneumoniae in 1 case.

Key words: MERS-CoV, diagnosis, respiratory agents, multiplex PCR, FilmArray

Background

The Middle East Respiratory Syndrome Coronavirus (MERS-CoV) is a novel coronavirus first described in the Arabian Peninsula in 2012.\textsuperscript{1} Humans may become infected through contact with contaminated respiratory secretions from infected patients or animals (mainly dromedaries). Although human-to-human transmission of the virus is low, MERS-CoV has been known to spread from ill people to others through close contact such as caring for or living with an infected person. Thus, transmission of MERS-CoV is more likely to occur in healthcare facilities,\textsuperscript{2} with most of the cases reported having been associated with nosocomial outbreaks.\textsuperscript{3} As of July 2017, 2040 laboratory-confirmed cases of MERS-CoV have been reported to the World Health Organization. To date, all cases of MERS have been linked to travel or residence in countries in and near the Arabian Peninsula. Moreover, exported cases\textsuperscript{4} and outbreaks\textsuperscript{5} outside the Arabian Peninsula have also been shown. The clinical syndrome ranges from asymptomatic and mild respiratory symptoms to severe acute respiratory disease and death, with a case fatality rate of \textsuperscript{-35\%}.\textsuperscript{4,5} To achieve early detection and avoid the transmission of the virus, suspicion of MERS-CoV should be raised whenever there are compatible clinical symptoms (acute febrile respiratory illness) and an epidemiological link (person resides or has travelled within a 14-day period before symptom initiation to the Middle East or any of the countries where MERS-CoV is known to be circulating in dromedary camels or where human infections have recently occurred).\textsuperscript{6} In addition, adequate measures of infection prevention and control should be taken until MERS-CoV infection is ruled out. Initial symptoms of MERS-CoV are non-specific and may resemble those of other respiratory agents. Having an accurate and early microbiological diagnosis without delaying MERS-CoV testing seems reasonable to improve patient management. In this study we evaluated the clinical and epidemiological aspects of the patients with suspicion of MERS-CoV attended in our centre and the definitive respiratory etiology identified through a comprehensive panel test.
Methods

We prospectively analysed all consecutive patients attending the Hospital Clinic of Barcelona with clinical and epidemiological suspicion of MERS-CoV according to the interim patient under investigation guidance from the CDC specifications and as established by the MERS-CoV suspected case management guidelines between June 2015 and January 2017. These guidelines specify that any patient with an epidemiologic risk and severe or milder respiratory illness should be considered as a patient under investigation for MERS-CoV. According to the protocol established by the Public Health Agency of Catalonia any clinical suspicion of MERS-CoV should be analysed in our hospital, which is the reference centre for MERS-CoV diagnosis in the region of Catalonia (North Eastern Spain) holding a laboratory with biosafety level II and providing rapid results 24 h a day, 7 days a week.

These patients remained in an isolation type III room until laboratory results became available. Respiratory samples (nasopharyngeal swabs and whenever possible a sputum) were collected and sent to the Microbiology laboratory. For the detection of MERS-CoV, RNA extraction of inactivated samples was performed using an automated system (Qiasympnomy; Qiagen, Hilden, Germany) followed by real-time reverse-transcription polymerase chain reaction (RT-PCR) targeting upstream of the 5′E gene, encoding protein E, as described previously. Synthetic RNA standards were used as the positive control. Additionally, all nasopharyngeal swabs samples were tested with a multiplex PCR system (FilmArray Respiratory Pannel, BioMérieux, Marcy l’Etoile, France). This is an integrated and automated system which allows the detection of 17 viral agents (influenza viruses A and B, parainfluenza viruses 1–4, respiratory syncytial virus, adenovirus, bocavirus, coronaviruses HKU1, NL63, 229E and OC43, metapneumovirus, human rhinovirus/enterovirus) and 3 bacterial agents (Bordetella pertussis, Chlamydia pneumoniae and Mycoplasma pneumoniae) in 1 h. All tests performed in the patients in this article were part of the routine clinical assessment at Hospital Clinic.

Results

A total of 14 patients fulfilled the definition of MERS-CoV suspected case during the study period. The median age of the patients was 30 years (interquartile range: 24–65), seven (50%) were males and seven (50%) were females. All the patients resided in the Middle East except for patient 13 who had been in Singapore when there was an alert for prevention of transmission of MERS-CoV due to the outbreak in the Republic of Korea. All patients presented respiratory signs or symptoms. Patients 1–11 presented milder respiratory illness (cough, odynophagia or dyspnea) and patients 12–14 presented pneumonia based on radiological evidence (Table 1). Real-time RT-PCR for the detection of MERS-CoV was performed in nasopharyngeal swab samples of all the patients and additionally from a sputum sample in patients 10, 12 and 14. All the samples tested negative for the detection of MERS-CoV. However, in 13 cases a respiratory pathogen was identified performing FilmArray multiplex PCR of the nasopharyngeal swabs. A respiratory virus was identified as the causative agent in 12

Table 1. Clinical and epidemiological characteristics of the patients included in the study and the identification of microorganisms using FilmArray Respiratory Pannel

<table>
<thead>
<tr>
<th>Patient</th>
<th>Age</th>
<th>Sex</th>
<th>City or country of origin</th>
<th>Clinical symptoms</th>
<th>Chest X-ray</th>
<th>Identification by FilmArray RP</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>24</td>
<td>M</td>
<td>Dammam (Saudi Arabia)</td>
<td>Chills, headache, myalgia, odynophagia and fever</td>
<td>NAD⁵</td>
<td>Parainfluenza virus 2</td>
</tr>
<tr>
<td>2</td>
<td>24</td>
<td>F</td>
<td>Bahrain</td>
<td>Odynophagia, dyspnoea, wheezing and fever</td>
<td>NAD⁵</td>
<td>Rhinovirus</td>
</tr>
<tr>
<td>3</td>
<td>20</td>
<td>F</td>
<td>Bahrain</td>
<td>Odynophagia and cough with purulent sputum</td>
<td>Not performed</td>
<td>Rhinovirus and Parainfluenza virus 4</td>
</tr>
<tr>
<td>4</td>
<td>65</td>
<td>F</td>
<td>Bahrain</td>
<td>Cough and odynophagia</td>
<td>Not performed</td>
<td>Rhinovirus and Parainfluenza Virus 4</td>
</tr>
<tr>
<td>5</td>
<td>68</td>
<td>F</td>
<td>Mecca (Saudi Arabia)</td>
<td>Cough without purulent sputum and fever</td>
<td>Not performed</td>
<td>Rhinovirus Influenza A H3</td>
</tr>
<tr>
<td>6</td>
<td>26</td>
<td>M</td>
<td>Al Bahah (Saudi Arabia)</td>
<td>Cough and dysthermia</td>
<td>Not performed</td>
<td>Rhinovirus</td>
</tr>
<tr>
<td>7</td>
<td>24</td>
<td>M</td>
<td>Riyadh (Saudi Arabia)</td>
<td>Cough, fever and odynophagia</td>
<td>NAD⁵</td>
<td>Rhinovirus</td>
</tr>
<tr>
<td>8</td>
<td>29</td>
<td>F</td>
<td>Jeddah (Saudi Arabia)</td>
<td>Cough and fever</td>
<td>Not performed</td>
<td>Influenza A H3</td>
</tr>
<tr>
<td>9</td>
<td>36</td>
<td>F</td>
<td>Riyadh (Saudi Arabia)</td>
<td>Cough and fever</td>
<td>Not performed</td>
<td>Influenza B</td>
</tr>
<tr>
<td>10</td>
<td>53</td>
<td>F</td>
<td>Iraq</td>
<td>Cough, fever and myalgia</td>
<td>Not performed</td>
<td>Negative</td>
</tr>
<tr>
<td>11</td>
<td>21</td>
<td>F</td>
<td>Kuwait</td>
<td>Cough and fever</td>
<td>Not performed</td>
<td>Influenza B</td>
</tr>
<tr>
<td>12</td>
<td>30</td>
<td>M</td>
<td>Kuwait</td>
<td>Non-productive cough, fever and vomiting</td>
<td>Infiltrate in the left lower lobe</td>
<td></td>
</tr>
<tr>
<td>13</td>
<td>75</td>
<td>M</td>
<td>Singapore</td>
<td>General malaise and disorientation</td>
<td>Infiltrate in the upper right lobe</td>
<td></td>
</tr>
<tr>
<td>14</td>
<td>76</td>
<td>M</td>
<td>Arabian countries</td>
<td>Cough, fever</td>
<td>Consolidation in the left lower lobe</td>
<td></td>
</tr>
</tbody>
</table>

⁵RP, respiratory panel.
⁶NAD, no abnormality detected.
cases and *M. pneumoniae* was identified as the causative agent in 1 case (Table 1).

**Discussion**

Although MERS-CoV does not seem to have an efficient human-to-human transmission, the high mortality rate of the virus and the fact that most of the cases have been related to healthcare-associated outbreaks reported in the Middle East and also in non-endemic areas from imported cases, makes necessary that potential new imported cases of MERS-CoV are early detected and isolated as soon as possible. Since the symptoms of MERS-CoV are non-specific and can be confused with infections due to other respiratory pathogens, the presence of other respiratory agents should also be investigated in patients with clinical suspicion of MERS-CoV. All the patients in the present study were negative for MERS-CoV, but the additional rapid testing for respiratory pathogens revealed the aetiological agent in most of the cases. This information allowed better patient management with rapid implementation of treatment and prevention and control measures.

Previous studies have shown that lower respiratory tract specimens contain higher viral loads and are more sensitive to virus detection, although upper respiratory tract specimens remain important for diagnosis and are more frequently available. The fact that all suspected cases were analysed in the reference laboratory underscores the importance of good quality sampling for respiratory tract infections with unusual location for a more sensitive approach.

Our results are in agreement with previous studies evaluating respiratory pathogens among pilgrims returning from an endemic area. However, our group of patients is heterogeneous not restricted to pilgrims and with patients returning from different endemic areas. Our study supports a comprehensive evaluation of respiratory agents in addition to MERS-CoV in patients with epidemiological and clinical suspicion of MERS-CoV infection coming to non-endemic areas.

**Conclusions**

Patients with respiratory symptoms returning from endemic MERS-CoV areas should be urgently screened for the possible presence of MERS-CoV. Additionally, it is of critical importance to early rule out other possible respiratory agents to avoid their transmission, prevent potential outbreaks and provide adequate patient management.

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**Author contributions**

Elsa Rubio: Study design, collection of microbiological and clinical data from the included patients and data analysis. Article writing. Miguel J. Martínez, Jordi Vila, Ma Àngues Marcos: Control and supervision of the microbiological procedures. Study design, article supervision and correction. Verónica Gonzalo, Josep Barrachina: Performance of the microbiological procedures. Núria Torner, Ana I Martínez, Mireia Jané, Anna Vililla, Antoni Trilla: Control and communication of the suspected cases. Article supervision and correction. Ana del Rio, Natalia Rodríguez-Valero, Maria Jesi Pinazo, José Muñoz, Alex Soriano: Patient management. Article supervision and correction.

**References**


