

Middle East Respiratory Syndrome Coronavirus and the One Health concept

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Middle East Respiratory Syndrome Coronavirus (MERS-CoV) is one of the major threats to the healthcare systems in some countries, especially in the Arabian Peninsula. MERS-CoV is considered an ideal example of the One Health concept. This is due to the animals especially dromedary camels play important roles in the transmission and sustainability of the virus, and the virus can be transmitted through aerosols of infected patients into the environment. However, there is some debate regarding the origin of MERS-CoV either from bats or other unknown reservoirs. The dromedary camel is the only identified animal reservoir to date. These animals play important roles in sustaining the virus in certain communities and may act as an amplifier of the virus by secreting it in their body fluids, especially in nasal and rectal discharges. MERS-CoV has been detected in the nasal and rectal secretions of infected camels, and MERS-CoV of this origin has full capacity to infect human airway epithelium in both in vitro and in vivo models. Other evidence confirms the direct transmission of MERS-CoV from camels to humans, though the role of camel meat and milk products has yet to be well studied. Human-to-human transmission is well documented through contact with an active infected patient or some silently infected persons. Furthermore, there are some significant risk factors of individuals in close contact with a positive MERS-CoV patient, including sleeping in the same patient room, removing patient waste (urine, stool, and sputum), and touching respiratory secretions from the index case. Outbreaks within family clusters have been reported, whereby some blood relative patients were infected through their wives in the same house were not infected. Some predisposing genetic factors favor MERS-CoV infection in some patients, which is worth investigating in the near future. The presence of other comorbidities may be another factor. Overall, there are many unknown/confirmed aspects of the virus/human/animal network. Here, the most recent advances in this context are discussed, and the possible reasons behind the emergence and sustainability of MERS-CoV in certain regions are presented. Identification of the exact mechanism of transmission of MERS-CoV from

camels to humans and searching for new reservoir/s are of high priority. This will reduce the shedding of the virus into the environment, and thus the risk of human infection can be mitigated.

1 **Middle East Respiratory Syndrome Coronavirus and the One Health Concept**

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17 **Running title:** MERS-CoV/human/animal interaction

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24 **Abstract:**

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26 healthcare systems in some countries, especially in the Arabian Peninsula. MERS-CoV is

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31 reservoirs. The dromedary camel is the only identified animal reservoir to date. These animals

32 play important roles in sustaining the virus in certain communities and may act as an amplifier of

33 the virus by secreting it in their body fluids, especially in nasal and rectal discharges. MERS-

34 CoV has been detected in the nasal and rectal secretions of infected camels, and MERS-CoV of

35 this origin has full capacity to infect human airway epithelium in both in vitro and in vivo

36 models. Other evidence confirms the direct transmission of MERS-CoV from camels to humans,

37 though the role of camel meat and milk products has yet to be well studied. Human-to-human

38 transmission is well documented through contact with an active infected patient or some silently

39 infected persons. Furthermore, there are some significant risk factors of individuals in close

40 contact with a positive MERS-CoV patient, including sleeping in the same patient room,

41 removing patient waste (urine, stool, and sputum), and touching respiratory secretions from the

42 index case. Outbreaks within family clusters have been reported, whereby some blood relative
43 patients were infected through their wives in the same house were not infected. Some
44 predisposing genetic factors favor MERS-CoV infection in some patients, which is worth
45 investigating in the near future. The presence of other comorbidities may be another factor.
46 Overall, there are many unknown/confirmed aspects of the virus/human/animal network. Here,
47 the most recent advances in this context are discussed, and the possible reasons behind the
48 emergence and sustainability of MERS-CoV in certain regions are presented. Identification of
49 the exact mechanism of transmission of MERS-CoV from camels to humans and searching for
50 new reservoir/s are of high priority. This will reduce the shedding of the virus into the
51 environment, and thus the risk of human infection can be mitigated.

52 **1. Introduction**

53 The main reason behind developing this article is to summarize the current understanding about
54 MERS-CoV in the context of the One Health concept. In this article, I highlight the known
55 information about the MERS-CoV infection and its pathogenesis in humans, the patterns of
56 MERS-CoV in dromedary camels, the potential roles of other animals in the transmission cycle
57 of MERS-CoV, and the interaction of MERS-CoV/humans/animals. I elaborate on how some
58 strategies can be used to stop or reduce the frequencies of MERS-CoV outbreaks based on the
59 One Health concept, identified some gaps in the literature, and drew conclusions. The one health
60 concept established to ensure the good health and well beings of the human, animal and the

61 environment. The human health is mainly affected by the environment health as well as the
62 animal health (Lerner & Berg 2015).

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64 **SURVEY METHODOLOGY**

65 For this review article, I conducted a literature search of the most up-to-date published articles on
66 MERS-CoV in the past 7 years. First, I focused the introduction section on the historical
67 background of coronaviruses and the One Health concept. Then, I highlighted the most up-to-
68 date literature from PubMed central, Google Scholar and ResearchGate on the interaction of
69 MERS-CoV/humans/animals. I identified some important gaps in the research dealing with
70 MERS-CoV/human/environment in the context of the One Health concept. I also summarized the
71 current acceptable theories on the emergence and evolution of MERS-CoV. Finally, I highlighted
72 progress to date in the control of MERS-CoV. Historically, MERS-CoV was first identified in
73 Saudi Arabia in a patient suffered from severe pneumonia and shortening of breath. The virus
74 was called the novel coronavirus at that time (Zaki et al. 2012). Another retrospective study
75 conducted in Jordan early 2012 revealed the detection of this novel coronavirus in 11 patients.
76 Eight out of them were from the health care workers (Hijawi et al. 2013).

77 **2. Coronaviruses: the past, present, and future**

78 Coronaviruses are a large group of viruses causing many health problems (respiratory,
79 enteric, and nervous syndromes) in various species of animals and humans. Six human
80 coronaviruses that have been identified to date (HCoV-229E, HCoV-OC43, HCoV-NL-63,

81 HCoV-HUK-1, SARS-CoV, and MERS-CoV). Two out of them emerged in the past 15 years
82 (Lau & Chan 2015), namely, the severe acute respiratory syndrome coronavirus (SARS-CoV)
83 and the Middle East Respiratory Syndrome Coronavirus (MERS-CoV). SARS-CoV emerged in
84 2003 in China and spread to many countries throughout the world (Peiris et al. 2003).
85 Approximately 8000 people were infected, and 10% of them died (Aronin & Sadigh 2004). Only
86 9 years later, MERS-CoV emerged in Saudi Arabia (Zaki et al. 2012). This is a relatively short
87 period for the emergence of a new coronavirus. One of the main reasons behind the rapid
88 emergence of new coronaviruses is the poor proofreading capability of their RNA polymerases
89 (Hofer 2013). This is in addition to the possibility of the recombination of different
90 coronaviruses (Makino et al. 1986), and it will not be surprising if new coronaviruses emerge in
91 the near future. MERS-CoV continues to pose great challenges to the healthcare system of some
92 countries in the Middle East and Arabian Peninsula. Since its discovery late in 2012 (Zaki et al.
93 2012), there are ongoing reports to the World Health Organization (WHO) from some countries
94 in the Middle East, especially the Arabian Peninsula, with spread to other countries around the
95 globe. According to the latest WHO statistics, there have been a total of 2,428 laboratory-
96 confirmed cases of MERS-CoV infection including at least 838 deaths (reported case fatality rate
97 of 35.0 %) (WHO 2018). The continuous ongoing reports on MERS-CoV suggesting the
98 presence of some factors favor its sustainability in certain regions. There are many uncertain
99 aspects of the virus evolution, pathogenesis, and transmission cycle. Unfortunately, recently,
100 there was some decline in the rate of research on the virus from different aspects (Hemida et al.
101 2017b). This hampered the production of new data about the MERS-CoV from different aspects.
102 Below, I summarize the current understanding of the virus in the context of the One Health
103 concept.

104 **3. Coronaviruses and the One Health concept**

105 The One Health concept is an interesting concept outlining the close interaction among humans,
106 animals and the environment (Destoumieux-Garzon et al. 2018). Currently, there are two
107 coronaviruses candidates representing the One Health Concept, SARS-CoV and MERS-CoV.
108 Animals play important roles in the transmission cycle of both viruses (Alshukairi et al. 2018;
109 Wang et al. 2005). Both viruses were proven to be of zoonotic origin (Gao et al. 2016). The palm
110 civet cat played important role in the transmission cycle of SARS-CoV (Wang et al. 2005). Some
111 patients proved to visit one restaurant serving the civet cats as a meal (Wang et al. 2005). Culling
112 of the civet cats resulted in marked decline in the reported SARS-CoV cases and now become
113 extinct. Many studies made a direct link between the exposure to camel and its meat and milk
114 products and MERS-CoV human cases (Reusken et al. 2014). Several studies reported the
115 presence of MERS-CoV specific antibodies in sera of human came in close contact with camels
116 (Reusken et al. 2014; Reusken et al. 2016). Meanwhile, MERS-CoV was isolated from air pf
117 positive dromedary camel herds in Saudi Arabia (Azhar et al. 2014).

118 **4. MERS-CoV in Humans**

119 MERS-CoV may infect a wide group of people ranging from very young ages, even infants
120 less than one year of age, to 109 years of age (CDC 2016). However, children are less likely to
121 be infected with MERS-CoV when compared to adults, and if infected, they tend to have
122 asymptomatic or mild disease (Arwady et al. 2016). The reason for this is still not entirely clear
123 and requires further study.

124 The case fatality rate is always very high in case of the immunocompromised infected
125 patients especially those who are suffering from chronic diseases such as cancer, diabetes, blood
126 pressure, kidney problems, etc. (Arwady et al. 2016).

127 Human-to-human transmission is reported in many cases. MERS-CoV replicates efficiently
128 in various *in vitro* and *ex vivo* models (Chan et al. 2014). Moreover, many family clusters and
129 hospital outbreaks were reported in the past 5 years (Arwady et al. 2016; Drosten et al. 2014;
130 Memish et al. 2013). This confirms the potential spread of MERS-CoV among those who are in
131 close contact in the population (Mollers et al. 2015). The most at-risk groups are healthcare
132 workers including nurses, medical doctors and other hospital staff and the elderly with
133 underlying chronic diseases (Arabi et al. 2014).

134 The prevalence rate of MERS-CoV in primary cases among males is relatively higher than
135 that of females (Darling et al. 2017), which may be because exposure to infected dromedary
136 camels is much higher in males than in females.

137 MERS-CoV infection triggers some unique interferons and cytokine gene expression
138 profiles. The virus seems to be a poor interferon inducer (Chan et al. 2014). This suggests the
139 potential immune evasion strategies triggered by the virus to hijack the host immune system and
140 may be responsible for the high fatality rate, at least in part. Viral spreading among people seems
141 to not yet be very efficient. Those in close contact are among the at-risk groups for infection
142 (Drosten et al. 2014), as observed in many hospital outbreaks as well as family clusters (Alfaraj
143 et al. 2018; Choi et al. 2017; Xiao et al. 2018). This suggests that transmission of the virus
144 among people requires exposure to a high viral load, which will sometimes produce active
145 infection in people who are in close contact. Several MERS-CoV family clusters have been

146 reported (Drosten et al. 2014). Interestingly, MERS-CoV is reported in the dromedary camels in
147 many African countries (Egypt, Nigeria, Tunisia, and Ethiopia), but no primary human cases
148 have been reported in these countries to date (Ali et al. 2017; Roess et al. 2016; van Doremalen
149 et al. 2017), which may be related to some variation in the circulating Asian and African strains
150 of MERS-CoV.

151 Some important deletions in the MERS-CoV currently circulating in dromedary camels from
152 Africa were recently reported (Chu et al. 2018). These deletions may explain at least in part the
153 reason behind the variations in the pathogenesis among the Asian and African strains of MERS-
154 CoV. Another potential reason behind the absence of human cases in the African countries is the
155 diverse cultural habits among people in Africa and the Arabian Peninsula (FAO 2016). People in
156 Arabian Peninsula get in more close touch with camels during the camel show, sports, trade than
157 in Africa. This make the human risk of exposure much higher in the AP than Africa. MERS-CoV
158 infection varies from severe respiratory illness accompanied by a high fever and respiratory
159 distress to mild asymptomatic cases. Patients are usually admitted to the intensive care unit
160 (ICU) and provided with a source of oxygen. Most cases result in pneumonia, which is fatal in
161 almost 40% of the affected patients (Hong et al. 2017; Rubio et al. 2018). Some patients may
162 develop renal failure [13]. Several MERS-CoV travel-associated infections were in many cases
163 associated with the Middle East (Bayrakdar et al. 2015; Rubio et al. 2018). Among these
164 reported was the Korean outbreak in early 2015 (Choi et al. 2017; Kim et al. 2017; Xiao et al.
165 2018). One Korean citizen visited some countries in the Middle East and then returned home ill.
166 This person visited several healthcare facilities in Korea. This resulted in the largest MERS-CoV
167 human outbreak outside the Arabian Peninsula (AP) (Xiao et al. 2018). This outbreak confirmed
168 the human-to-human transmission. During this outbreak, MERS-CoV was isolated from air

169 samples from the hallways of the healthcare facilities close to the hospitalized patients (Xiao et
170 al. 2018). This at least explains in part the rapid development of MERS-CoV hospital outbreaks.

171 **5. MERS-CoV in animals**

172 Since the discovery of MERS-CoV late in 2012 (Zaki et al. 2012), many research groups
173 have searched for its potential animal reservoir/s. Dromedary camels are the only currently
174 proven reservoir for MERS-CoV (Hemida et al. 2014; Hemida et al. 2017b; Reusken et al. 2014;
175 Reusken et al. 2016). Interestingly, others were able to trace the virus back 30 years ago in
176 dromedary camel specimens in retrospective studies (Corman et al. 2014; Hemida et al. 2014;
177 Reusken et al. 2014). All these data suggest that the virus has been circulating for decades
178 without being recognized. Although the actual and typical clinical features of the MERS-CoV
179 natural infection in dromedary camels is not well documented to date, very few studies reported
180 these patterns under experimental infection approaches (Adney et al. 2014). Based on these
181 findings, camels do not show any pathognomonic signs despite a subtle fever and mild nasal
182 discharge for up to 6 days post-infection (dpi) (Hemida et al. 2014). Meanwhile, shedding of the
183 infectious virus was reported in the experimentally infected camels started at 2 dpi up to the 7th
184 dpi (Adney et al. 2014). Interestingly, viral RNA was still detected at 35 dpi (Adney et al. 2014),
185 though it is not known whether the viral RNAs may act as potential sources of infection similar
186 to some other positive-sense RNA viruses. No viral shedding in the oral secretions, rectal swabs,
187 urine, or sera of these animals has been reported (Adney et al. 2014), in contrast to the detection
188 of the virus in the fecal specimens and swabs under natural viral field infection (Hemida et al.
189 2014). These finding suggesting differential patterns of MERS-CoV infection between natural
190 and experimental approaches. Further studies are required to understand the natural MERS-CoV
191 infection in dromedary camels, which may be achieved by conducting long-term longitudinal

192 studies as well as careful monitoring of the virus infection in large populations of camels. On
193 necropsy examination of MERS-CoV, experimentally infected dromedary camels revealed only
194 mild-to-moderate inflammatory reactions in the upper respiratory tract (Khalafalla et al. 2015).
195 Detection of the viral antigens in the tissue sections of the turbinate bone and the upper
196 respiratory tract was reported (Adney et al. 2014). Interestingly, seroconversion of the inoculated
197 animals was reported to begin at 14 dpi (Hemida et al. 2014), indicating that MERS-CoV
198 induces a robust humoral immune response after infection. More recently, one longitudinal study
199 reported the possibility of MERS-CoV infection in seropositive animals. This raises concern
200 about the role of the antibodies in the protection of the MERS-CoV infection (Hemida et al.
201 2017a). It seems that all the members of the family *Camelidae* (dromedary, alpaca, and llamas)
202 are susceptible to MERS-CoV infection (Corman et al. 2014; Vergara-Alert et al. 2017). David
203 et al. (2018) reported the presence of antibodies against MERS-CoV in some alpacas and llamas
204 in Israel but only used commercial ELISA kits, and they did not address the possibility of cross-
205 reactivity with other coronaviruses especially BCoV. It had been previously showed that there is
206 clear cross-reactivity between MERS-CoV and BCoV in dromedary camels (David et al. 2018).
207 Interestingly, one study showed an absence of any detectable antibodies of MERS-CoV in the
208 sera of Bactrian camels (Chan et al. 2015), though this is the only study to report this finding
209 concerning the seronegativity of Bactrian camels to MERS-CoV. It is not well known whether
210 the absence of detectable MERS-CoV antibodies in the sera of Bactrians camels is due to the
211 geographical location of the tested animals in Mongolia, far from the Middle East and Africa.
212 This may be supported by similar findings in dromedary camels in Australia and the Canary
213 Islands (Crameri et al. 2015). Another possibility is that this might be due to some genetic
214 factors, which contribute to the resistance of Bactrians to MERS-CoV infections; this point is

215 worthy of further investigation. Experimental MERS-CoV infection in both alpacas and llamas
216 showed a similar pattern to that of dromedary camels (Crameri et al. 2016; Vergara-Alert et al.
217 2017), which suggested that both animals might act as a model animal for the study of MERS-
218 CoV *in vivo*. However, the experimental infection of pigs with MERS-CoV did not reveal as
219 much infection as that reported in alpacas and llamas (Vergara-Alert et al. 2017). Active MERS-
220 CoV particles were neither retrieved from the experimentally infected animals nor from the close
221 contact non-infected animals during the duration of this study (Vergara-Alert et al. 2017). This
222 result suggested that pigs might not play an active role in the transmission of MERS-CoV.
223 Although bats are considered the main reservoir for many coronaviruses, their roles in the
224 MERS-CoV still need further clarifications. One study reported the presence of MERS-CoV in
225 one specimen collected from bats in Saudi Arabia (Memish et al. 2013). The genome sequence
226 of this particular virus showed almost 100% identity to a MERS-CoV index case (Memish et al.
227 2013). More recently, Jamaican fruit bats were found to be permissible for MERS-CoV infection
228 (Munster et al. 2016). However, MERS-CoV-infected bats did not show any apparent clinical
229 signs; however, viral shedding was reported in the swabs from bats up to 9 dpi. The clinical
230 profiles and viral shedding curve during the course of the MERS-CoV infection in these bats
231 look similar to that of dromedary camels (Munster et al. 2016), yet the amount of infectious viral
232 shedding in bats is less compared to that in dromedary camels. This species of bat is not the most
233 relevant for MERS-CoV infection, but this study offers some insights into the molecular
234 pathogenesis of MERS-CoV in bats. Interestingly, another study revealed the expression of
235 MERS-CoV receptors (dipeptidyl peptidase-4, DPP4) in the respiratory and digestive tracts of
236 some insectivorous bats (Widagdo et al. 2017). An interesting study tested the potential roles of
237 other species in the transmission of MERS-CoV such as cattle, sheep, goats, donkeys, buffaloes,

238 mules, and horses from Egypt, Tunisia and Senegal (Kandeil et al. 2019). This study revealed the
239 presence of neutralizing antibodies in the sera of some sheep and goats. Meanwhile, viral RNA
240 was detected in swabs collected from some sheep, goats and donkeys from these countries
241 (Kandeil et al. 2019). Several attempts were made to identify an appropriate experimental animal
242 model for MERS-CoV. The Syrian hamster was non-permissive to MERS-CoV infection (de Wit
243 et al. 2013). Experimental infection of this animal neither develops any clinical signs or
244 pathology nor produces any cytokines after infection (de Wit et al. 2013). This was in contrast to
245 New Zealand white rabbits, which showed active infection after inoculation with the MERS-
246 CoV (Monchatre-Leroy et al. 2017). Furthermore, both rhesus macaques and common
247 marmosets supported MERS-CoV infection (Yu et al. 2017). Additionally, both the transgenic
248 and the transduced mice expressing the dipeptidyl peptidase 4 human receptors worked as a
249 model for MERS-CoV studies (Zhao et al. 2015). Interestingly, a new study reported the
250 seropositivity of some sheep and goat to MERS-CoV from Tunisia, Senegal and Egypt (Kandeil
251 et al. 2019). Same study reported the detection of the viral RNAs in samples from cow, sheep,
252 goat and donkeys from Egypt (Kandeil et al. 2019). This highlights the importance of continuous
253 surveillance and searching for new reservoir/s for the MERS-CoV transmission cycle.

254 **6. MERS/human/animal interaction**

255 It is now well accepted that human exposure to MERS-CoV-infected dromedary camels is a
256 predisposing factor to human infection, particularly in immunocompromised people (Zumla et al.
257 2015). Based on the latest WHO reports, the prognosis of MERS-CoV infection is poor for
258 elderly patients who have chronic diseases such as cancer, diabetes, kidney failure, etc. (Arabi et
259 al. 2014). Transmission of MERS-CoV from dromedary camels to humans has been proven
260 indirectly in some previous reports (Azhar et al. 2014). One study showed strong evidence of

261 direct transmission of MERS-CoV from an infected camel to its owner, which was confirmed by
262 comparing the viral genome sequencing of the virus isolated from the infected dromedary camel
263 to that isolated from its owner. Both viruses were almost a 100% match (Azhar et al. 2014).
264 Meanwhile, this study reported the detection of MERS-CoV nucleic acid in air samples from the
265 indicated dromedary camel barn during the active course of the viral infection (Azhar et al.
266 2014).

267 There is a debate about the role of the dromedary camel's milk and meat products and by-
268 products in the transmission of MERS-CoV. Experimental introduction of MERS-CoV to raw
269 milk revealed little difference between the virus stock in milk and that kept in DMEM (van
270 Doremalen et al. 2014). Due to their culture, some people in the Middle East would drink raw
271 camel milk in efforts to seek treatment for some diseases such as diabetes. The authors
272 acknowledge that MERS-CoV was introduced into the dromedary camel milk at a high dose and
273 that the viral RNA was detected in a limited concentration in the camel's milk (van Doremalen et
274 al. 2017). Thus, drinking the raw camel milk poses a great risk to the people consuming this milk
275 without any heat treatment or pasteurization (van Doremalen et al. 2014; Zhou et al. 2017). One
276 study connected the infection of some people to the drinking of the milk from one infected camel
277 (Memish et al. 2015). However, another study was conducted in Qatar to assess the possibility of
278 acquiring the infection from contaminated teats and udders of infected she-camels during the
279 process of milking (Reusken et al. 2014), though no active MERS-CoV shedding in milk has yet
280 to be reported. Further studies are encouraged to come to a conclusion about the potential role of
281 raw camel milk in the transmission of MERS-CoV. Nonetheless, the role of camel meat in the
282 transmission of MERS-CoV has not been studied carefully to date. Thus, special attention should
283 be paid to the efficient cooking of camel meat and its products as well as thorough boiling of

284 camel milk. It is suggested that people not drink raw camel milk to avoid any risk of infection
285 not only with MERS-CoV but also with other pathogens such as Brucellosis (Garcell et al.
286 2016). Some studies reported that MERS-CoV is one of the occupational zoonotic viral diseases,
287 as was claimed based on some studies investigating the seroconversion of some at-risk group of
288 people to MERS-CoV. This study reported the presence of specific MERS-CoV antibodies in
289 approximately 3 % of the workers in some slaughterhouses in Qatar (Farag et al. 2015). On the
290 other hand, some studies reported the absence of any detectable antibodies in the sera of some
291 herdsmen, veterinarians, and slaughterhouses in Saudi Arabia (Hemida et al. 2015). One possible
292 explanation for the variations between the two studies is the difference in the sensitivity of the
293 techniques used. Those two studies used two different techniques to report the presence/absence
294 of the MERS-CoV antibodies in the sera of at-risk people (Farag et al. 2015; Hemida et al.
295 2015). Regardless, further investigation on a large-scale basis is required to solidify this
296 conclusion about MERS-CoV.

297 **7. Gaps on the MERS-CoV-related research**

298 There is more to be known about the molecular biology of MERS-CoV. Identification of the
299 DPP-4 as viral receptors does not rule out the presence of other co-receptors or
300 transcription/translation factors that favor the virus infection in a certain host. There are many
301 immune evasion strategies triggered by MERS-CoV to hijack the host immune responses, and
302 the mechanisms of such strategies have not been well studied. Moreover, there are many
303 unknown aspects especially in the context of the MERS-CoV/human/animal interaction.
304 Meanwhile, some studies were conducted on a small scale or with low numbers of
305 animals/specimens and reported some important conclusion. These studies need further
306 confirmation, and refinement of some of these observations is urgently needed in the near future.

307 Here, we highlight some gaps in the research regarding the evolution and transmission of MERS-
308 CoV. Presumably, there might be an unidentified reservoir in the transmission cycle of MERS-
309 CoV. Although respiratory infection still is the main route of MERS-CoV infection, the exact
310 mechanism of transmission of MERS-CoV from dromedary camels to humans is still not well
311 understood. The possibility of another reservoir in the transmission cycle of MERS-CoV has not
312 been ruled out. Thus, there might be a missing link in the chain of the human/camel interaction.
313 Meanwhile, the exact modes of transmission of MERS-CoV from dromedary camels to humans
314 have not been well clarified, and the typical pattern of the natural MERS-CoV infection in
315 dromedary camels has not been well studied. Additionally, the potential role of most camel
316 secretions and excretions has not been fully understood. The seroprevalence of MERS-CoV was
317 reported in the dromedary camels from different countries in Africa and Asia (Ali et al. 2017;
318 Hemida et al. 2014), though feral camels in Australia and the Canary islands were found to be
319 seronegative (Crameri et al. 2015). The reason behind this phenomenon may be due to these
320 regions being isolated lands and away from the MENA region as described above; it may also be
321 due to the absence of an active camel movement between the Middle East and Africa and these
322 regions of the world. Very few studies reported the cross-reactivity between MERS-CoV and
323 other coronaviruses such as the bovine coronavirus (BCoV) that might infect dromedary camels.
324 It is unknown if this might be the reason behind the high seroprevalence of MERS-CoV among
325 the dromedary camel population. This may be due to the high frequency of exposure to the
326 MERS-CoV infection during the camel's life, the cross-reactivity of other coronaviruses, or an
327 unknown mechanism related to the dromedary camel's immune system. These considerations
328 require further studies. There is ongoing demand for the development of novel diagnostic assays
329 for coronaviruses, and special interest should be paid to those techniques that enable the

330 simultaneous detection of the viral nucleic acids and those that can simultaneously distinguish
331 between the antibodies for several coronaviruses. Furthermore, it is not well understood why
332 only the Bactrian camels among the family Camelidae did not seroconvert to MERS-CoV
333 infection (Chan et al. 2015). The genetic susceptibility of certain human populations, especially
334 blood-related people, is not clear in the context of MERS-CoV infection. There are several levels
335 of human exposure to dromedary camels, such as camel attendants, workers in camel abattoirs,
336 veterinarians inspecting their carcasses, and camel owners. Those groups of people are in close
337 contact with camels for various amounts of time and are considered to be a high-risk group of
338 people due to the long time they spend in close contact with the dromedary camels. Meanwhile,
339 there is an urgent need to develop a risk scoring system for human exposure to the dromedary
340 camels.

341 **8. Current theories in the MERS-CoV/human/animals interaction**

342 It is believed that there is some unidentified reservoirs in the context of MERS-CoV
343 transmission presenting the virus to the community. This virus is able to infect dromedary
344 camels, which act as an amplifier host for the virus, favoring the circulation of the virus in some
345 camel herds. The virus has the ability to circulate among the animals in the same herd and the
346 camel herds in close proximity to them (Figure 1). MERS-CoV in camels has the full potential to
347 infect the human especially immunocompromised persons. Once the virus infects a human, there
348 is always a possibility of infecting other people, especially closely-related individuals (figure 1),
349 including household relatives and workers plus healthcare workers such as doctors and nurses.
350 Infection depends on the level of exposure to the infected person, and MERS-CoV infection in
351 humans ranges from very severe cases of pneumonia to death. Currently available data indicate
352 that severely infected individuals can shed the infectious virus into the environment (Kim et al.

353 2016), though there are few data regarding the capacity of mildly infected individuals to transmit
354 the virus. Asymptomatic individuals, however, are unlikely to transmit the virus (Moon & Son
355 2017).

356 **9. Potential reasons for the emergence and the spread of MERS-CoV**

357 There are many factors behind the emergence, sustainability and spread of MERS-CoV. The
358 presence of an unidentified MERS-CoV reservoir in the transmission cycle is still considered,
359 and this unknown reservoir may contribute substantially to the suitability of the virus in certain
360 regions. Dromedary camels remain the amplifier of the virus; the close contact of these animals
361 to the human population in certain regions of Africa and Asia may pose a great risk for human
362 infection and indirectly contribute to the spread of the virus. Additionally, public animal markets,
363 especially for dromedary camels, may act as an amplifier of the virus. This poses a great risk to
364 the surrounding community. Interesting study addressed the mapping of MERS-CoV cases in
365 association with the environmental conditions and camel exposure (Reeves et al. 2015). This
366 study revealed that camel exposure is a key predisposing factor for some of MERS-CoV human
367 cases (Reeves et al. 2015). The lack of active surveillance programs for respiratory viruses,
368 especially coronaviruses, may result in many subclinical or mild cases of MERS-CoV being
369 missed in a certain population. These patients may shed the virus in their secretions and may act
370 as a source of infection to other persons in close contact with them. Although many MERS-CoV
371 vaccine and drug candidates are being researched, none are available yet. All these factors may
372 favor the sustainability of MERS-CoV in certain regions.

373 **10. Current progress on the control of MERS-CoV**

374 Interestingly, the case fatality rate of MERS-CoV among the affected population dropped from
375 almost 50% in 2012 to 34 % early 2019 (WHO 2018), and we may relate this progress in the
376 control of MERS-CoV over the past 7 years to many factors. First, identification of the main
377 reservoir of the virus, namely, the dromedary camel (Hemida et al. 2014). Second, continuous
378 molecular and serological surveillance of MERS-CoV among the dromedary camel population in
379 the Arabian Peninsula and Africa (Corman et al. 2014; Farag et al. 2015; Hemida et al. 2017a;
380 Hemida et al. 2017b; Khalafalla et al. 2015; Reusken et al. 2014). Currently, testing the
381 population of camels in regional camel markets is associated with shutting down of the market in
382 case of positive shedding of MERS-CoV by the animals. I believe this will substantially
383 minimize the risk of community-acquired infections through these positive populations. Third,
384 vaccination of dromedary camels, especially animals under two years of age, will have a great
385 impact on the reduction of the viral shedding from these animals to the surrounding community.
386 This will also have a great positive impact on the reduction of the number of reported human
387 infections. Fourth, there has been progress in the current understanding of viral tropism,
388 pathogenesis, and mode of transmission in the past five years (Chan et al. 2014; Widagdo et al.
389 2017). Fifth, new strategies have been adopted to reduce the spread of infection in health care
390 units (Rajakaruna et al. 2017). Sixth, some therapeutic and control approaches for MERS-CoV
391 such as cyclosporine, ribavirin and interferon show promising trends for the treatment of MERS-
392 CoV-infected patients (Al-Tawfiq et al. 2014; de Wilde et al. 2013). Meanwhile, good progress
393 has been made in screening large numbers of drugs/therapies for the treatment of MERS-CoV
394 (Han et al. 2018; He et al. 2019; Niu et al. 2018; Totura & Bavari 2019). This may lead to the
395 development of some effective novel drugs against MERS-CoV infection in the near future.

396 **11. One Health-based interventions to stop MERS-CoV outbreaks**

397 To stop MERS-CoV outbreaks, there are several strategies to be adopted in the context of the
398 One Health concept. Some strategies are related to the animal, while others are related to human
399 health. The main objective is to minimize or stop the viral shedding from dromedary camels to
400 the environment (Figure 2). This may be achieved in many ways including regular monitoring of
401 the population of dromedary camels. Active animal shedders need to be identified, and
402 quarantine measures should applied until they stop shedding the virus. Vaccination of young
403 dromedary camel calves should occur during their first 6 months of life, which will minimize the
404 chances of these animals becoming infected and actively passing the virus to older animals and
405 then to the environment. Reorganization and reshaping of the camel industry includes allocating
406 the camel markets away from the cities. Global awareness concerning the necessity of thorough
407 boiling and cooking of the camel milk and meat products, respectively, should take place.
408 Animal abattoirs should be established far away from large cities. They should not use mixed-
409 animal platforms, and each platform should deal with one species of animal. Thorough
410 decontamination of animals' biological wastes in abattoirs should occur using the appropriate
411 standard protocols. Regular surveillance of MERS-CoV among the population especially during
412 the active peak of virus shedding by the animals should occur during November to April every
413 year. People who are in close contact with the camels should wear proper personal protective
414 equipment at all times.

415 **12. Conclusions**

416 At almost 7 years after its emergence, there are ongoing reports of MERS-CoV infection from
417 time to time. This may be related to many unknown aspects of the viral evolution and pathogenesis.
418 Some of these unknown aspects are the following. (1) Little is still known about virus/host
419 interactions and how MERS-CoV hijacks the host immune system. (2) Potential reservoirs in the

420 context of the MERS-CoV/human/animal network need to be identified. (3) Why do MERS-CoV-
421 infected dromedary camels not exhibit obvious clinical signs during active viral shedding? (4)
422 Does the presence of neutralizing antibodies in the sera of animals protect them against future
423 active infection by the virus? (5) Does vaccination of dromedary camels have a positive impact on
424 controlling the spread of MERS-CoV among dromedary camels? More research is urgently needed
425 to explore the unknown aspects of the MERS-CoV/human/animal network.

426 **Figure legends**

427

428 **Figure 1: Current acceptable theories in the MERS-CoV/human/animals interaction**

429 There might be an unknown reservoir in the transmission cycle of MERS-CoV. Bats play a
430 role in the context of MERS-CoV transmission. The virus is transmitted to dromedary camels
431 through an unknown mechanism. The dromedary camels act as amplifying hosts for the virus.
432 MERS-CoV is transmitted from dromedary camels to humans through the respiratory aerosols and
433 some other unknown mechanisms. The virus is then transmitted among the human population
434 through respiratory routes. The human-to-human transmission has been confirmed. The human-
435 to-camel transmission still needs further clarification. Question marks indicate the non-confirmed
436 phenomenon.

437

438 **Figure 2. Some interventions based on the One Health-based to stop MERS-CoV outbreaks**

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667

Figure 1

Current theories regarding the MERS-CoV/human/animal interaction

There might be an unknown reservoir in the transmission cycle of MERS-CoV. Bats play a role in the context of MERS-CoV transmission. The virus is transmitted to dromedary camels through an unknown mechanism. The dromedary camels act as amplifying hosts for the virus. MERS-CoV is transmitted from dromedary camels to humans through the respiratory aerosols and some other unknown mechanisms. The virus is then transmitted among the human population through respiratory routes. The human-to-human transmission has been confirmed. The human-to-camel transmission still needs further clarification. Question marks indicate the non-confirmed phenomenon.

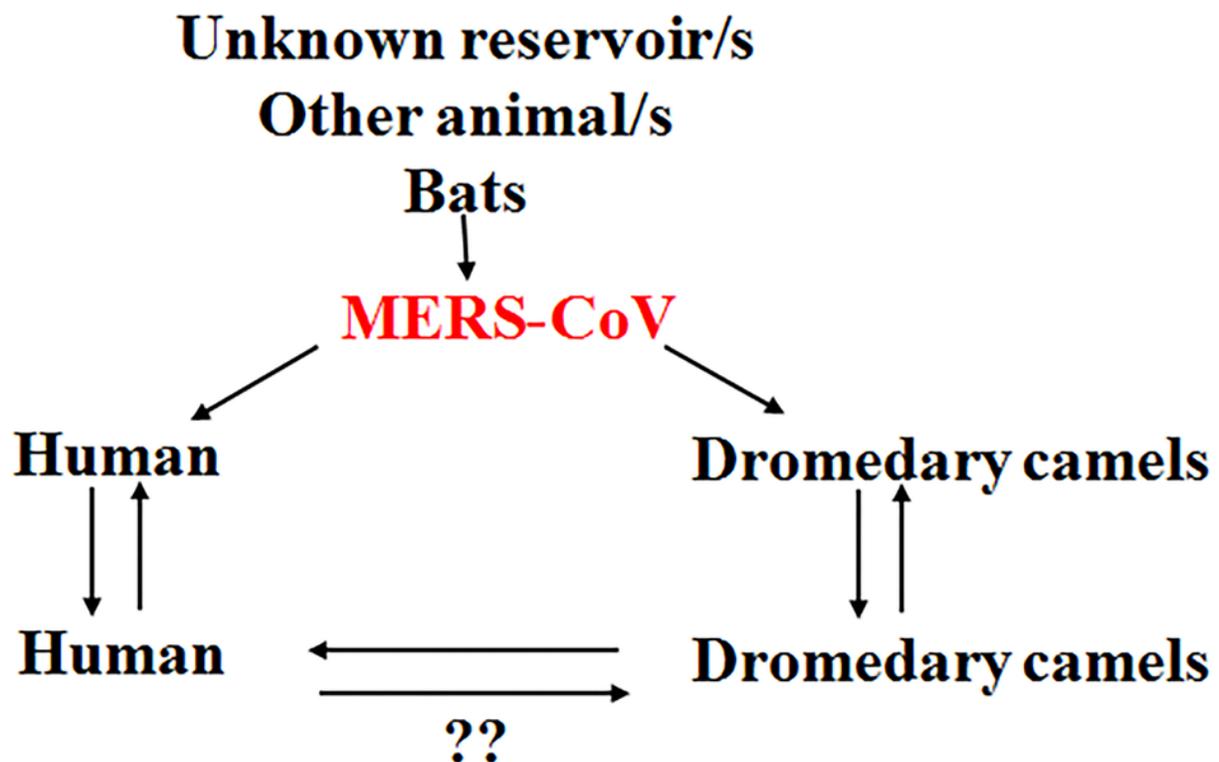


Figure 2

Figure 2

Some interventions based on the One Health-based to stop MERS-CoV outbreaks

