

Middle East respiratory syndrome

Ziad A Memish, Stanley Perlman, Maria D Van Kerkhove, Alimuddin Zumla



The Middle East respiratory syndrome coronavirus (MERS-CoV) is a lethal zoonotic pathogen that was first identified in humans in Saudi Arabia and Jordan in 2012. Intermittent sporadic cases, community clusters, and nosocomial outbreaks of MERS-CoV continue to occur. Between April 2012 and December 2019, 2499 laboratory-confirmed cases of MERS-CoV infection, including 858 deaths (34·3% mortality) were reported from 27 countries to WHO, the majority of which were reported by Saudi Arabia (2106 cases, 780 deaths). Large outbreaks of human-to-human transmission have occurred, the largest in Riyadh and Jeddah in 2014 and in South Korea in 2015. MERS-CoV remains a high-threat pathogen identified by WHO as a priority pathogen because it causes severe disease that has a high mortality rate, epidemic potential, and no medical countermeasures. This Seminar provides an update on the current knowledge and perspectives on MERS epidemiology, virology, mode of transmission, pathogenesis, diagnosis, clinical features, management, infection control, development of new therapeutics and vaccines, and highlights unanswered questions and priorities for research, improved management, and prevention.

Introduction

During the past 17 years three lethal zoonotic diseases of humans caused by novel coronaviruses, severe acute respiratory syndrome (SARS) in November, 2002,¹ Middle East respiratory syndrome (MERS) in April, 2012,² and more recently coronavirus disease (COVID-19) in December, 2019.³ All three causative coronaviruses, respectively, SARS-CoV, MERS-CoV, and SARS-related coronavirus-2 (SARS-CoV-2) have focused the attention of global public health authorities and are listed in the WHO Blueprint list for priority pathogens⁴ because of their pandemic potential, threat to global health security, and the absence of any effective treatments.

SARS-CoV was first identified in humans in Guangdong, China, in November, 2002 and subsequently spread rapidly worldwide to 29 countries, resulting in 8098 human SARS cases with 774 deaths (9·6% mortality).^{1,5} The SARS epidemic ended abruptly in July, 2003,¹ and no human cases of SARS have been detected over the past 15 years. MERS-CoV was first identified as causing human disease when it was isolated from a lung sample of an adult patient who was admitted at a hospital in Jeddah, Saudi Arabia, with severe pneumonia and died of multiorgan failure.⁶ A retrospective study then linked MERS-CoV to a hospital outbreak in April, 2012, in Jordan.⁷ MERS-CoV is considered a zoonotic pathogen, with MERS-CoV-infected dromedary camels being the animal source of infection to humans.^{8,9}

Unlike SARS-CoV, which was contained within a year of emerging, MERS-CoV continues to circulate and cause human disease with intermittent sporadic cases, community clusters, and nosocomial outbreaks in the Middle East with considerable risk of spreading globally.^{2,10} Several outbreaks of human-to-human MERS-CoV transmission have occurred, the largest outside the Middle East occurring in South Korea in 2015.^{11,12} This outbreak was associated with substantial morbidity and mortality, as well as having substantial economic, social, and health security effects.¹²

SARS-CoV-2 (initially named 2019-nCoV) was detected in December 2019 after sequencing of clinical samples

from a cluster of patients in Wuhan, China who developed pneumonia of unknown cause.¹³ This outbreak is rapidly evolving and as of March 3, 2020 there have been 88 948 laboratory confirmed cases of COVID-19, with 3043 deaths (3·4% mortality). Outside China, there have been 8774 cases reported from 64 countries with 128 deaths.³

This Seminar reviews current knowledge and provides an update on MERS-CoV epidemiology, virology, mode of

Search strategy and selection criteria

We searched publications in the English language literature on PubMed and Google Scholar (dates we used to refine our search: Sept 1, 2012, to Dec 31, 2019). We used the search terms "Middle East Respiratory Syndrome" OR "MERS-CoV" OR "MERS" in combination with the terms "Coronavirus", OR "Middle East", OR "Epidemiology", "Aetiology", "Dromedaries", "Camels", "Bats", OR "Transmission", OR "Virology" OR "Pathogenesis", "Clinical features" OR "Infection control", "Diagnostic tests", OR "Diagnosis" OR "Treatment", OR "Management", OR "drugs", "Therapies", OR "Prevention" OR "Vaccines" OR "Nosocomial" OR "Hospital". We also searched websites of global and national public health agencies such as WHO, WHO-Regional Office for the Eastern Mediterranean, UN Food and Agriculture Organization, US Centers for Disease Control and Prevention (CDC), Public Health England (UK), World Organisation for Animal Health, European Centre for Disease Prevention and Control (ECDC), and the Saudi Ministry of Health Middle East respiratory syndrome (MERS) portal. We reviewed publications on MERS from WHO, CDC, ECDC, Center for Infectious Disease Research and Policy (University of Minnesota, USA), and ProMED (Program for Monitoring Emerging Diseases) websites. We focused our review on publications published in the past 36 months to update the *Lancet* Seminar on MERS published in September, 2015. We also selected reference lists and substantive reviews of articles identified by this search strategy that we judged relevant to provide readers with additional details and more references than this Seminar can accommodate.

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College of Medicine, Alfaisal University, Riyadh, Saudi Arabia (Prof Z A Memish FRCP); Research Center, King Saud Medical City Ministry of Health, Riyadh, Saudi Arabia (Prof Z A Memish); Hubert Department of Global Health, Rollins School of Public Health, Emory University, Atlanta, GA, USA (Prof Z A Memish); Department of Microbiology and Immunology, and Department of Pediatrics, University of Iowa, Iowa City, IA, USA (Prof S Perlman PhD); Infectious Hazards Management, Health Emergencies Programme, World Health Organization, Geneva, Switzerland (M D Van Kerkhove PhD); Department of Infection, Division of Infection and Immunity, Centre for Clinical Microbiology, University College London, London, UK (Prof Sir A Zumla FRCP); and National Institute for Health Research Biomedical Research Centre, University College London Hospitals, London, UK (Prof Sir A Zumla)

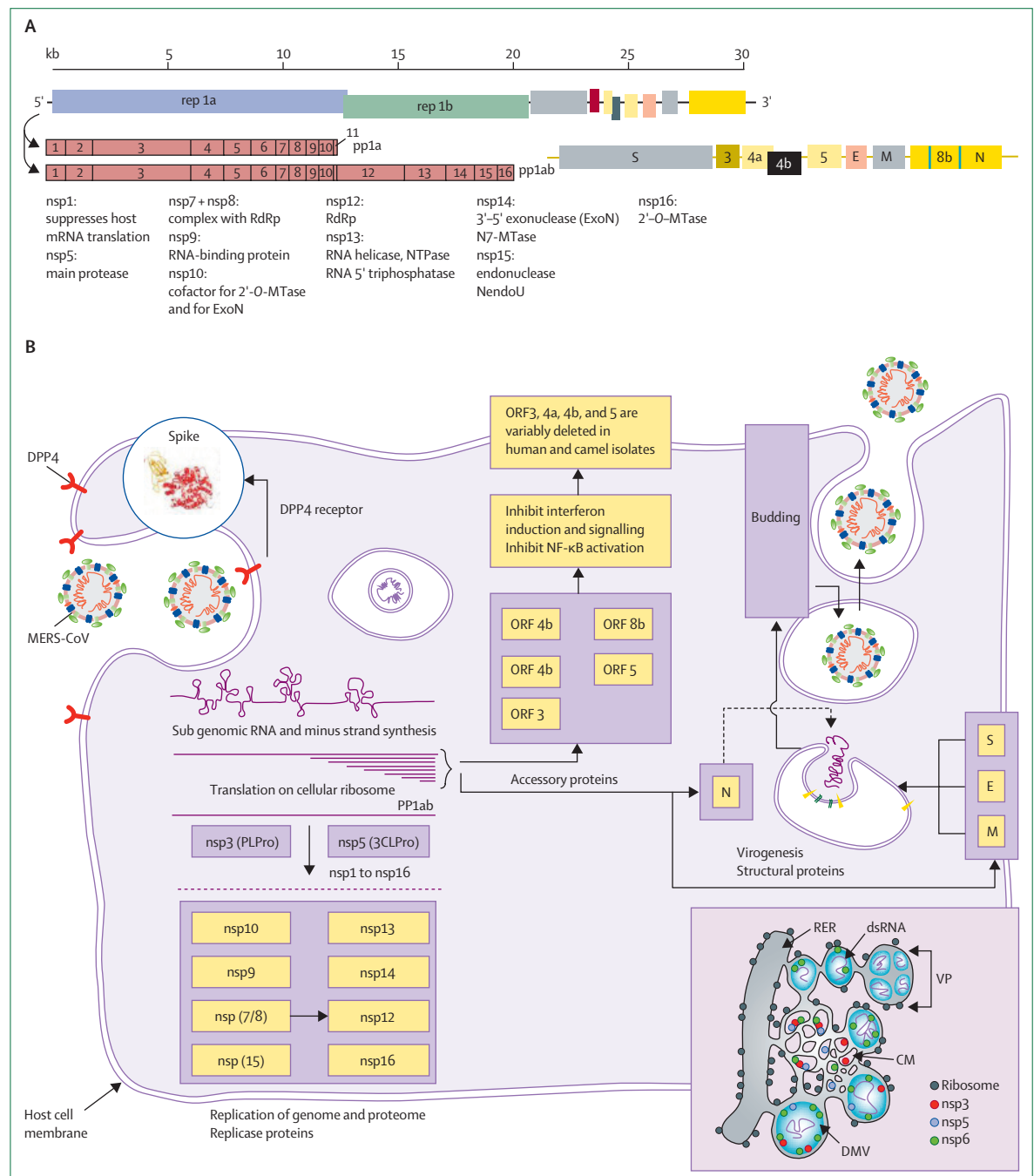
Correspondence to: Prof Sir Alimuddin Zumla, Department of Infection, Division of Infection and Immunity, Centre for Clinical Microbiology, University College London, London, NW3 0PE, UK a.zumla@ucl.ac.uk

transmission, pathogenesis, diagnosis, clinical features, management, infection control, and development of new therapeutics and vaccines, and also highlights unanswered questions and priorities for research, improved management, and prevention.

Virology

MERS-CoV, SARS-CoV, and SARS-CoV-2, are members of the *Coronaviridae* family of the order *Nidovirales*. Other human coronaviruses generally cause mild respiratory

infections (eg, HCoV-229E, HCoV-NL63, HCoV-OC43, and HCoV-HKU1). MERS-CoV, like SARS-CoV and SARS-CoV-2, can cause highly lethal disease in humans. MERS-CoV is a large single-strand positive-sense RNA virus (figure 1).^{14–18} The 30–31 kb coronavirus genome encodes a large number of proteins, which might confer versatility in adapting to new environments and enhance cross-species transmission. MERS-CoV has four structural proteins: spike (S) protein, envelope (E) protein, membrane (M) protein, and nucleocapsid (N) protein.



The S protein is a type I transmembrane glycoprotein located as a trimer on the virus surface and consists of S1 and S2 subunits. It has crucial roles in binding, fusion, and entry into host cells. The S1 subunit has a receptor binding domain that binds to the host cellular receptor dipeptidyl peptidase 4 (DPP4). The S2 subunit has two regions, heptad repeats 1 and 2 (HR1 and HR2), which rearrange to form a six-helix bundle to enable membrane fusion. The E and M proteins are present in the viral membranes and are required for viral assembly, budding, and intracellular trafficking.

MERS-CoV enters host cells via binding of its S protein to the host cell DPP4 receptor (figure 1; appendix pp 2–3).^{15,17} DPP4 is less abundant in the nasal cavity and epithelial cells of the upper airways but is highly expressed on the epithelial cells of the distal airways and type I and II pneumocytes in the lung alveoli, non-ciliated bronchial epithelial cells, endothelial cells, and some haemopoietic cells.¹⁵ DPP4 is also widely expressed on the epithelial cells of several other organs and tissues such as the kidneys, intestine, liver, thymus, and bone marrow—thus widespread dissemination of MERS-CoV could occur in the body. The structures of the S protein of MERS-CoV and other coronaviruses have been determined using cryo-electron microscopy^{16,17} and provide

the basis for rational design of protective antibodies and other therapeutics.

Epidemiology

A primary MERS-CoV infection is defined by WHO as a laboratory-confirmed MERS-CoV infection that has no direct epidemiological link to a human MERS-CoV infection, and was acquired outside of a health-care facility presumably from direct or indirect contact with the reservoir host—dromedary camels.¹⁹

A secondary MERS-CoV infection is defined by WHO as a laboratory-confirmed MERS-CoV infection with a direct epidemiological link to an individual with confirmed or probable MERS-CoV infection.¹⁹

See Online for appendix

Geographical distribution

Laboratory-confirmed MERS-CoV human infections¹⁹ are reported to WHO as a requirement under the International Health Regulations (2005). The geographical distribution of countries reporting laboratory-confirmed human MERS cases and the numbers reported over time are shown in figures 2 and 3. Between April, 2012 and end of December, 2019, 2499 laboratory-confirmed human cases of MERS-CoV infection, including 858 deaths (34·3% mortality) were reported from 27 countries in all continents (figure 2) to WHO, the majority of which were reported by Saudi Arabia (2106 cases, 780 deaths).

Sources of infection

The origin of all primary human MERS-CoV infections remains unknown. Dromedary camels are a host reservoir species for the MERS-CoV (appendix pp 3–5).^{20–23} Humans can acquire MERS-CoV through direct or indirect contact with infected dromedary camels or infected patients^{19–30} (figure 4A). However not all cases infected in the community report contact with dromedary camels (appendix p 4–5). The transmission of MERS-CoV from dromedary camels to humans is now well documented in the Arabian Peninsula,^{23–25} but the extent to which this transmission is occurring in countries outside the Arabian Peninsula requires definition. Experimental infections have shown that infected dromedaries shed MERS-CoV from their nasal secretions with minimal signs of illness, which is limited to rhinorrhoea.^{27–29} Full-genome phylogenetic analysis indicates that evolution of diverse MERS-CoV lineages in camels have caused human infections, perpetuating a low barrier for interspecies transmission.^{21,29,30} However, African MERS-CoV lineages in camels imported into Saudi Arabia have not established themselves in camels in Saudi Arabia,^{9,30} indicating potential differences in transmission dynamics and selection pressure. Analysis of population dynamics shows that Arabian viruses can maintain endemicity without introduction of additional lineages.^{9,30} WHO is supporting molecular and serological studies at the dromedary–human interface in several countries in

Figure 1: MERS-CoV structure, genome organisation, and replication

MERS-CoV encodes a large replicase-transcriptase polyprotein (rep1A and rep1B), which is processed into 16 non-structural proteins (nsp). These proteins are required for the formation of the replicase-transcription complex, for cleavage of the polyprotein, and for immune evasion. Structural proteins (spike [S], envelope [E], membrane [M], and nucleocapsid [N]) and accessory proteins (ORF3, 4a, 4b, 5, 8b) are encoded in the end third of the 3' end of the genome of the genome. MERS-CoV binds to its cellular receptor DPP4 via the S protein, which is processed by host proteases to expose a fusion peptide. The viral genome is then released into the cytoplasm, where it is translated on host ribosomes into rep1A and rep1B proteins. The polyprotein is cleaved by two viral-encoded proteases, encoded by nsp3 and nsp5. Proteins involved in genome and subgenome replication and transcription include nsp12 (the RNA-dependent RNA polymerase [RdRP]) and two associated proteins, nsp7 and nsp8.¹²⁷ Nsp13 (a helicase), nsp14 (which encodes an N7 methyltransferase and ExoN, a protein required for genome fidelity), and nsp16 (a 2'-O methyltransferase) are also necessary for optimal genomic and subgenomic RNA synthesis. MERS-CoV transcription involves the synthesis of subgenomic RNAs, which encode the structural and accessory proteins located at the 3' end of the genome. Subgenomic and genomic RNAs are co-terminal, sharing the same 5' leader and 3' sequences. Subgenomic RNAs code for structural and accessory proteins (ORF3, 4a, 4b, 5, and 8b). ORF8b is encoded within the N gene (marked with purple lines). These accessory proteins are believed to have immunoevasive properties, but are not essential for replication, and are variably deleted in human and camel virus isolates.^{128–130} Genomic replication occurs on membrane structures such as double membrane vesicles (DMVs), convoluted membranes (CM), and vesicle packets (VP), which are merged DMV³¹ that have been formed from the rough endoplasmic reticulum (RER) by the combined action of nsp3, nsp4, and nsp6 (lower right). After synthesis on replicase-transcription complexes, RNA is encapsidated by the N protein and transported to the ERGIC (endoplasmic reticulum–Golgi compartment), where budding into membranes containing the S, E, and M proteins occurs before release from the cell. 3CLPro=chymotrypsin-like protease. ExoN=exonuclease. MERS-CoV=Middle East respiratory syndrome coronavirus. MTase=methyltransferase. NendoU=nidoviral uridylylate-specific endoribonuclease. NF-κB=nuclear factor κ-light-chain-enhancer of activated B cells. NTPase=nucleoside-triphosphatase. pp1a=polyprotein 1a. PLPro=papain-like protease.

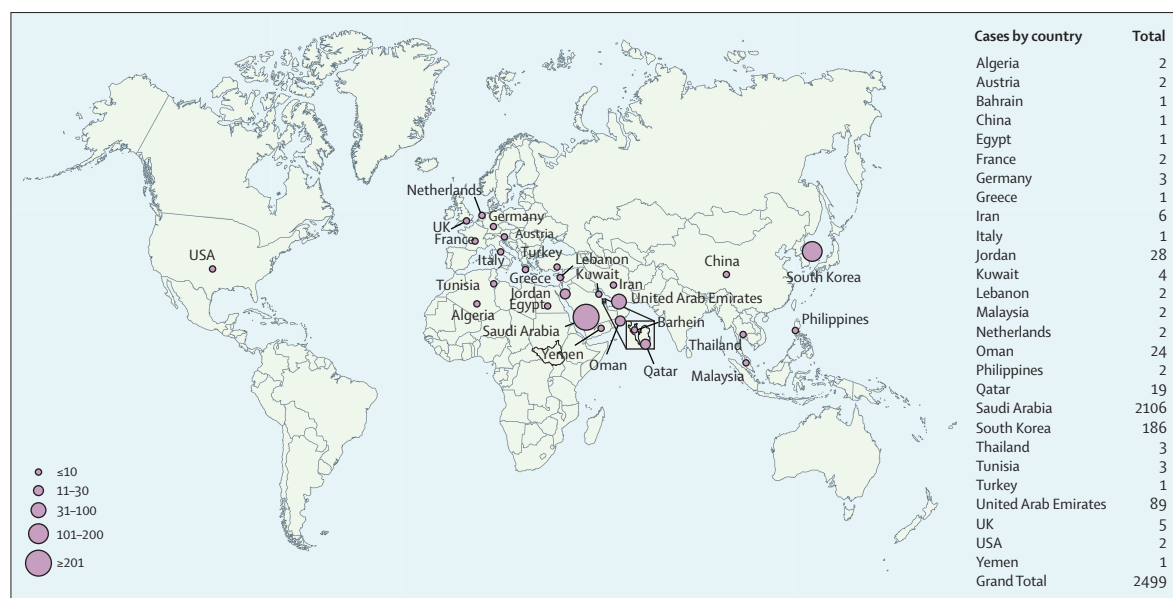


Figure 2: Geographical distribution of reported human infections of MERS-CoV

Cases reported up to and including Dec 31, 2019. MERS-CoV=Middle East respiratory syndrome coronavirus. Reproduced from Dr Mamunur R Malik, WHO Regional Office for the Eastern Mediterranean.

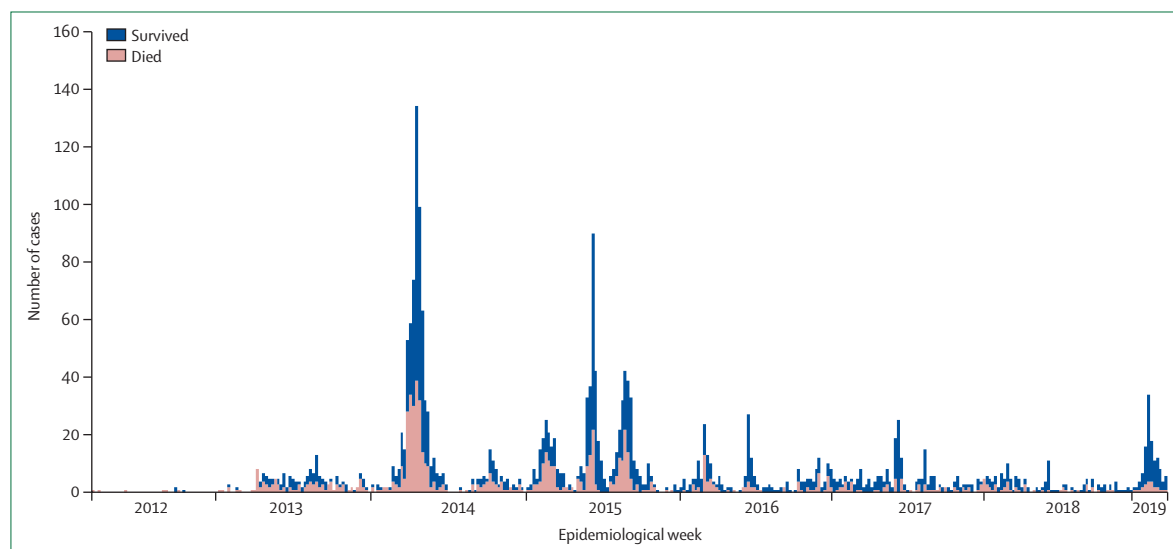


Figure 3: Global MERS reported to WHO by week, 2012–19

Reproduced from WHO website, by permission of Dr Mamunur R Malik, WHO Regional Office for the Eastern Mediterranean. MERS=Middle East respiratory syndrome.

Africa and south Asia.³¹ The phylogenetic relationship of complete genomes of MERS-CoV strains obtained from camels and humans has been published.³²

Epidemiological patterns of human cases

The epidemiological patterns of MERS-CoV in humans have remained consistent since MERS-CoV was first identified in 2012.² Primary cases often report direct or indirect contact with dromedary camels and present across a wide clinical spectrum from mild to severe fulminant disease. Individuals with severe primary

MERS-CoV infections are often older than 65 years with comorbidities, and symptoms can present late. Individuals with mild primary MERS-CoV infections are often missed by current surveillance systems since they usually do not present to health-care facilities.

Secondary cases have resulted from human-to-human transmission among close contacts. To date, secondary transmission has occasionally occurred between close contacts of individuals with laboratory-confirmed MERS-CoV in household settings.^{33,34} However, secondary transmission in health-care facilities has repeatedly

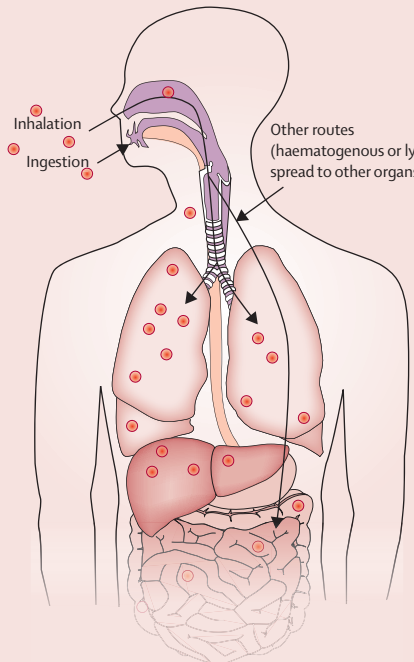
A Sources of MERS-CoV	B Probable MERS-CoV routes of entry, circulation, and organ involvement	C Clinical, imaging, and laboratory features	
<p>Animal and environmental sources</p> <p>MERS-CoV-infected camels</p> <ul style="list-style-type: none"> Camel nasal secretions Camel excreta (urine, saliva and faeces) Camelid birth products (amniotic fluid, fetal membranes and placenta) Camel food products (milk, meat) Other unknown? <p>Human sources</p> <p>Symptomatic and subclinical MERS-positive individuals in:</p> <ul style="list-style-type: none"> The community Family households Family compounds Hostels Health-care facilities and hospitals Camel farms <p>Up to 25% report direct or indirect contact with camels and approximately 50% are the result of human-to-human transmission</p>	<p>MERS-CoV host cell DPP4 receptor found in:</p> <ul style="list-style-type: none"> bronchial epithelium, lung parenchyma, interstitium (endothelial cells), kidneys, intestines, liver, thymus, haemopoietic cells (leukocytes, macrophages, dendritic cells, mononuclear lymphoid cells) 	<p>Presenting symptoms</p> <p>General</p> <ul style="list-style-type: none"> Fever (>38°C) Chills or rigours Lethargy Anorexia Malaise, lethargy Myalgia Body aches <p>CNS</p> <ul style="list-style-type: none"> Headache Confusion <p>Upper respiratory tract</p> <ul style="list-style-type: none"> Runny nose Sneezing Sore or tickly throat <p>Lower respiratory tract</p> <ul style="list-style-type: none"> Cough (dry or productive with sputum) Shortness of breath Chest pain Haemoptysis <p>Gastrointestinal tract</p> <ul style="list-style-type: none"> Decreased appetite Nausea Vomiting Diarrhoea Abdominal pain Abdominal discomfort 	<p>Incubation period</p> <ul style="list-style-type: none"> Mean: 5·2 days (95% CI 1·9–14·7) Range 2–14 days <p>Sex distribution</p> <ul style="list-style-type: none"> Males 64%, females 36% <p>Abnormal investigations</p> <ul style="list-style-type: none"> Imaging: chest x-ray and CT changes Leucopenia Lymphopenia Thrombocytopenia Elevated LDH Elevated ALT Elevated AST <p>Complications</p> <ul style="list-style-type: none"> Severe acute respiratory syndrome Respiratory failure Liver failure Renal failure Multiorgan failure Septic shock <p>Factors associated with increased mortality</p> <p>Comorbidities:</p> <ul style="list-style-type: none"> Diabetes Chronic lung disease (COPD, asthma) Chronic kidney disease Chronic liver disease Chronic heart disease Malignancies Immunosuppressive drugs Age >65 years Presence of pleural effusion Low serum albumin <p>Case fatality rate</p> <ul style="list-style-type: none"> Overall (global): 36·0% Saudi Arabia: 41·8% (22%–69%) South Korea: 20·4% (14·5–47%)

Figure 4: Epidemiological, clinical, and laboratory features of MERS

ALT=alanine aminotransferase. AST=aspartate aminotransferase. MERS-CoV=Middle East respiratory syndrome coronavirus.

occurred in several countries and has, on occasion, resulted in large outbreaks (appendix p 5)—such as Jeddah, Saudi Arabia (2014), Seoul, South Korea (2015), and Riyadh, Saudi Arabia (2015, 2016, and 2018). In health-care settings, human-to-human transmission occurs between patients, between patients and health-care workers, and from patients to visitors.^{21,33} Approximately half of the MERS-CoV cases reported to WHO² to date have resulted from human-to-human transmission in health-care facilities.^{21,33}

Risk factors and transmission

Known risk factors for MERS-CoV acquisition, transmission, and outbreaks are noted (panel 1). MERS patient data reported to WHO includes information on exposures and known risk factors during the 14 days before symptom onset, or during the 14 days before laboratory confirmation was reported (in the case of asymptomatic infection). Exposure data include travel history to MERS-CoV endemic countries, direct or indirect contact with dromedary camels or their products, contact with humans with MERS-CoV infection, and visits to health-care facilities containing patients infected with MERS-CoV are risk factors for acquiring MERS-CoV

infection (appendix pp 4–5).^{2,24,25,31–45} MERS-CoV has been detected in camel products (eg, raw milk, meat, blood, urine, or birth products); however, genomic studies to definitively show transmission from these products to humans have not yet been done. Human primary intestinal epithelial cells, small intestine explants, and the intestinal tract are highly susceptible to MERS-CoV and can sustain viral replication. After infection of mice transgenic for the expression of the host DPP4 receptor, histological examination showed MERS-CoV enteric infection in all inoculated mice with progression to infection of the lung tissues, indicating the development of sequential respiratory infection.³⁹

Transmission via contact with contaminated hospital environments is possible during outbreaks, although genomic studies providing an evidence base are lacking. Since environmental contamination is a potential source of infection, it is prudent to take precautionary infection control measures.

Nosocomial transmission

Health-care-associated outbreaks of MERS-CoV affecting inpatients, health-care workers, and visitors are characteristic of MERS (panel 2) and account for approximately

Panel 1: Risk factors for MERS-CoV infection, transmission, and outbreaks^{21,34,38,132–138}**Community risk factors**

- Indirect or direct contact with MERS-CoV-infected camels, camel nasal secretions, milk, urine, faeces, meat, or birthing products
- Travel to the Middle East and contact with camels or attendance at a health facility
- Exposure to, or contact with, patients with MERS-CoV in the community, or at healthcare facilities
- Underlying chronic medical conditions

Nosocomial risk factors for human-to-human transmission and outbreak amplification

- Late diagnosis of MERS-CoV infection
- Lack of awareness in health-care workers of the possibility of MERS in patients with a fever
- Overcrowded or contaminated health-care facilities, especially emergency departments, dialysis units, and crowded inpatient wards
- Exposure of hospital staff or other patients to symptomatic patients with MERS
- Delayed implementation of infection control measures
- Poor compliance with MERS-specific infection control guidelines (hand hygiene, droplet and contact precautions, environmental cleaning)
- Poor compliance with appropriate personal protective equipment when assessing patients with febrile respiratory illness
- Aerosol-generating procedures or invasive procedures in patients with MERS (eg, nebulisers, resuscitation, intubation, and ventilation)
- Lack of proper isolation room facilities
- Distance between patient beds of less than 1 m
- Friends and family members staying as caregivers in overcrowded health-care facilities

MERS=Middle East respiratory syndrome. MERS-CoV=Middle East respiratory syndrome coronavirus.

50% of reported cases.^{2,20,21,33,40–45} Large nosocomial outbreaks have occurred in Abu Dhabi in United Arab Emirates, Seoul in South Korea and several cities in Saudi Arabia (appendix p 5). The 2013 outbreak in Al Hasa, Saudi Arabia, involved 23 people in three different hospitals and was linked to poor infection control practices, aerosol-generating procedures, continuous positive pressure ventilation, and cardiopulmonary resuscitation.⁴² In June, 2017, an outbreak of 34 MERS-CoV cases (including 17 in health-care workers) occurred in a hospital in Riyadh, Saudi Arabia,⁴³ where the primary case was a 47-year-old man admitted for emergency intubation. Before a diagnosis of MERS, this individual had contact with 220 health-care workers, patients, and visitors. Another 2017 outbreak in Riyadh involved three hospitals, with 44 MERS cases in patients, health-care workers, and family members. The outbreak arose from three severely ill people who were diagnosed with MERS late in the disease course, and involved in a superspreading event.⁴⁴ In 2015, South Korea had the largest MERS-CoV outbreak outside the Middle East with 186 cases reported, resulting from a single imported case from an individual who had returned from travelling in the Middle East. The outbreak included 185 secondary cases who acquired infection in

16 health-care facilities and 80% of transmission events were attributed to five superspreading events.⁴⁵

Pathogenesis and pathology

The mechanisms underlying the pathogenesis of MERS-CoV remain to be defined because autopsies are generally not done either for religious and cultural reasons or to prevent environmental contamination with subsequent infection of health-care workers. Therefore, there are few data on the histopathological changes in patients with MERS-CoV infection, even in severe disease.^{46,47} The only two autopsies available show that viral infection was confined predominantly to the respiratory tract, although viral particles were detected in the kidney in one of the reports.⁴⁷ Further, ex vivo human kidney cultures were shown to support productive MERS-CoV infection.⁴⁸ However, it is not known whether infection of this organ contributes to worse outcomes compared with infections confined to the respiratory tract. Viral RNA, but not infectious MERS-CoV, was detected in the blood for at least 2 weeks after diagnosis. Whether this viral RNA represented an extrapulmonary infection is not clear, since MERS-CoV neutralising antibody was detected in the serum at the same time in some patients.⁴⁹

Pathological changes in the lungs include evidence of focal haemorrhagic necrotising pneumonia with exudative diffuse alveolar damage, indistinguishable from findings detected in severe pneumonia caused by other viral agents.

Several experimentally infected animal models for MERS have been developed (appendix p 6).^{50,51} Methods for gene silencing and editing are being further developed, therefore it is likely that additional viral and host factors important in MERS-CoV pathogenesis will be identified.

Protective and deleterious immune responses

MERS-CoV pathogenesis reflects a balance between coronavirus-induced protective and pathogenic host immune responses and direct cytotoxic effects of the virus.⁵² According to studies of patients infected with SARS-CoV and of mice and other animals infected experimentally with MERS-CoV, successful resolution of MERS and long-term protection from re-infection likely requires well coordinated innate and adaptive B-cell and T-cell responses. In patients with SARS, innate immune responses characterised by an extended period of cytokine secretion (eg, IFN α and IFN β) was correlated with delayed antibody responses and poor management outcomes; this prolonged innate response has not yet been shown in patients infected with MERS-CoV.⁵³ MERS-CoV productively infects activated human T cells and induces delayed cytokine responses after infection of human myeloid cells in vitro, both of which could impair virus clearance and the development of an effective immune response.^{54–56}

During the acute phase of MERS, robust virus-specific CD8 T-cell responses were detected in most patients with severe or moderate disease, with antibody and CD4 T-cell responses appearing later in the disease course.⁵⁷ T-cell and antibody responses were reliably detected 2–3 weeks after diagnosis, although they were detected earlier in some patients.^{49,57} Studies of MERS survivors showed that MERS-CoV-specific antibody responses tended to be lower and transient in patients with mild or subclinical disease when compared with patients with severe disease, in whom MERS-CoV-specific antibody responses were detected for at least 2 years.^{57–60} In contrast, T-cell responses were detected in all MERS survivors for at least 2 years.⁵⁹

The transitory nature of the antibody response in mild disease and greater stability of the T-cell response suggest that induction of both will be required for optimal long-term protection; the measurement of both will enhance the accuracy of prevalence studies.

Clinical features

Incubation period

The specific exposures that lead to sporadic MERS-CoV infections are unknown, therefore it is impossible to estimate the incubation period in primary cases. However, based on data from cases of human-to-human transmission, the incubation period is a median of 5–7 days, with a range of 2–14 days (median 5.2 days [95% CI 1.9–14.7]).^{2,10,21,61–63} Immunocompromised patients can present with longer incubation periods of up to 20 days.⁶⁴

Presentation in adults

The clinical presentation of patients infected with MERS-CoV ranges from asymptomatic or mild upper respiratory illness to rapidly progressive pneumonitis, respiratory failure, acute respiratory distress syndrome, septic shock, and multiorgan failure with fatal outcome (figure 5).^{2,10,11,61–70} It is not possible to distinguish patients with a legitimate asymptomatic MERS-CoV infection from those who develop mild disease because reports to WHO provide signs and symptoms information only at the time of reporting. Many individuals remain asymptomatic whereas some go on to develop mild disease, which is why WHO classifies these individuals as mild or asymptomatic. Asymptomatic-to-mild infection rates of 25–50% have been reported.

The signs and symptoms associated with MERS are non-specific, with or without multisystem involvement, and thus could be mistaken for other causes of respiratory tract or gastrointestinal illnesses.^{2,10,11,61–70} Therefore, the clinical diagnosis of MERS can be easily missed. Patients with MERS can typically present with fever, chills, rigors, headache, a non-productive cough, sore throat, arthralgia, and myalgia followed by dyspnoea. Other associated symptoms include coryza, nausea, vomiting, dizziness, sputum production, diarrhoea, and abdominal pain. Some patients with MERS can present

Panel 2: MERS prevention and interventions^{21,31,33,36,67}

The pathogen: MERS-CoV

- Anti-viral therapeutics including host-directed therapies or immune-based therapies are in development.

Intermediate host reservoir: dromedary camels

- Vaccinating dromedary camels, the source for primary human infection, is a proposed approach to preventing human infection.
- A recombinant camelpox virus expressing the S protein has been shown to induce neutralising antibodies in camels.
- As compared to older camels, juvenile camels have higher rates of viral shedding. Past infection is not protective.

Humans with contact with infected camels

- Basic hand hygiene practices.
- Gloves and face masks when handling infected camels and camel products.
- Pasteurising camel milk.
- Avoid ingesting raw camel meat, dairy products, or urine.
- Vaccine development for people with high camel exposure.

Human cases

- High clinical awareness of the possibility of MERS-CoV infection and making early diagnosis.
- Isolate suspected cases early and institute infection control measures.
- Face masks and hand hygiene measures for all HCW and family contacts.
- Occupationally exposed people have a higher risk of infection compared to the general public. Human surveillance is focused on diseased people, and those with subclinical or asymptomatic infection and could serve as a source of infection of a susceptible person with comorbidities.

Nosocomial and community outbreaks

- Nosocomial cases and outbreaks are diminishing due to institution of stringent infection control measures but still account for 35–50% of cases.
- More than 20 human vaccines have been developed and are being tested in animals—some are in human phase 1 or 2 trials.

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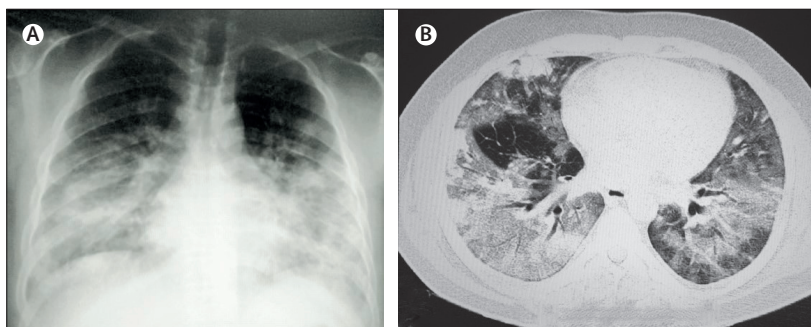


Figure 5: Chest imaging abnormalities in patients with Middle East respiratory syndrome
(A) Chest x-rays showing bilateral extensive diffuse and focal opacities. (B) Chest CT scan showing bilateral extensive ground-glass reticulonodular shadowing with bronchiolar wall thickening.

with atypical symptoms of mild respiratory illness without a fever and a gastrointestinal illness that precedes the development of pneumonia.^{64,67,68} Neuromuscular manifestations include hypersomnolence, weakness, and

tingling in the extremities similar to Guillain-Barré syndrome or virus-related sensory neuropathy.⁶⁸ Co-infection of MERS-CoV with other respiratory viruses (such as parainfluenza virus, rhinovirus, influenza A or B virus, respiratory syncytial virus, enteroviruses, and human metapneumovirus) and nosocomial bacterial infections has been reported in patients receiving intensive care.^{61,67}

Immunocompromised patients or those with comorbidities

Immunocompromised patients or patients with chronic heart, liver, and kidney disease can present with atypical features, such as a longer incubation period, a longer period from initial PCR positivity to symptom onset, persistent prolonged viral shedding, and increased mortality.^{64–70} MERS-CoV causes more severe disease in people older than 60 years, and those with chronic comorbid diseases such as renal disease, cancer, chronic lung disease, immunosuppression, and diabetes.

Children, pregnant women, and returning travelers

In children younger than 5 years, MERS-CoV infection appears to be rare, and usually presents as mild disease with cough as the predominant clinical symptom. Acute and severe respiratory illness (ie, a fatal outcome after multiorgan failure) has been noted in only two of 38 reported paediatric cases and these were in boys aged 9 months and 2 years with comorbidities (infantile nephrotic syndrome and cystic fibrosis, respectively).^{71–74} Data on the prevalence of MERS-CoV in pregnant women are limited to case reports, and the clinical presentations and mortality rates are similar to non-pregnant women.⁷⁵

MERS should be kept on the differential diagnosis list for ill travellers returning home from MERS-CoV endemic areas who present with acute febrile illness (temperature $\geq 38^{\circ}\text{C}$), aches and pains, headache, cough, dyspnoea, diarrhoea, nausea, vomiting, unexplained leucopenia (white blood cell count $< 3.5 \times 10^9/\text{L}$), and thrombocytopenia (platelets $< 150 \times 10^9/\text{L}$)—either with or without prominent respiratory symptoms.^{2,10,11,61–70} It is important that clinicians are alert to the possibility that patients could have MERS in all health-care settings where the virus is circulating so that an accurate diagnosis can be made and infection control measures implemented efficiently.⁷⁶

Clinical specimen types for diagnosis

The type and quality of the clinical specimen are important in the laboratory diagnosis of MERS-CoV infection. Both upper respiratory tract specimens (oropharyngeal or nasopharyngeal swabs) and lower respiratory tract specimens (sputum, endotracheal aspirate or lavage) should be analysed whenever possible.^{77–79} Patients with MERS might not shed the virus during the early stage of their illness; thus, initial negative results from upper respiratory samples

do not rule out the possibility of MERS and patients should be retested using a lower respiratory tract sample. Several studies of MERS-CoV viral load measurements show that upper respiratory tract specimens have lower viral loads than lower respiratory specimens.^{11,12,79,80} MERS-CoV RNA has been detected in blood, urine, and stool—but at much lower viral loads than in the respiratory tract.^{81,82}

Clinical specimens must be collected by trained staff using appropriate personal protective equipment—eg, when taking nasopharyngeal and oropharyngeal specimens, Dacron or rayon swabs specifically designed for collecting specimens for virology must be used.^{77,78,83} These swab kits should contain virus transport medium. A single negative test result does not exclude the diagnosis and repeat sampling and testing is strongly recommended.⁸³ A minimum of two samples, preferably from the lower respiratory tract, are needed to exclude MERS-CoV after initial assessment. To confirm clearance of the virus, respiratory samples should be collected sequentially (every 2 to 4 days) over ensuing days until there are two consecutive negative results at least 24 h apart in people who are clinically recovered.⁷⁷

Laboratory testing

WHO guidelines for testing should be followed.⁷⁷ MERS-CoV testing must be performed in appropriately equipped biosafety laboratories by staff trained in the relevant technical and safety procedures. National or WHO guidelines on the laboratory biosafety should be followed in all circumstances.⁸³

Three real-time RT-PCR (rRT-PCR) assays for routine detection of MERS-CoV have been developed targeting upstream of the E protein gene (upE) and open reading frame (ORF)1b and ORF1a. The assay for the upE target is considered highly sensitive and is recommended for screening, with the ORF1a assay (ORF1a assay is considered more sensitive than ORF1b assay).⁷⁷

An updated roadmap for MERS-CoV product development lists all available diagnostics and other diagnostics in the developmental phase.⁸⁴ Several diagnostic tests are under development to accelerate turnaround times from sample analysis to result.^{85–87} A 2018 rapid and specific assay for the detection of MERS-CoV combines the reverse transcription loop-mediated isothermal amplification technique and a vertical flow visualisation strip (RT-LAMP-VF) to detect the nucleocapsid gene of MERS-CoV.⁸⁶ The RT-LAMP-VF assay provides detection of MERS-CoV within 35 min and is easier to perform than the WHO-recommended rRT-PCR method.⁸⁶

Several serological assays are available for detection of MERS-CoV: ELISA, recombinant spike immunofluorescent assay, and spike pseudoparticle neutralisation and microneutralisation assay.^{87–90} A case confirmed by serology requires evidence of seroconversion in two samples, ideally taken at least 14 days apart, by a screening (eg, ELISA, immunofluorescence assay) and a neutralisation assay.^{77,78}

Imaging

A range of abnormal but non-specific chest x-ray findings are seen in patients with MERS.^{91,92} These abnormalities include unilateral or bilateral bronchovascular shadowing, interstitial infiltrates, reticular opacities, reticulonodular shadowing, nodules, pleural effusions, and patchy to confluent consolidation (appendix pp 18–19). Lower lobes tend to be affected more than upper lobes early in the course of MERS and rapid opacification of lungs and progression to acute respiratory distress syndrome can occur.

High-resolution CT might show ground glass opacities (figure 5B) in early CT findings (with or without consolidation) followed by interlobular septal and intralobular interstitial thickening with peripheral and lower lobe involvement^{93,94} within the first week of MERS-CoV infection. During subsequent weeks, a so-called crazy-paving lung abnormality, cavitation, tree-in-bud pattern and centrilobular nodules, constrictive obliterative bronchiolitis, bronchiolitis obliterans, peribronchiolar air-trapping, thickened peripheral bronchioles, and organising pneumonia have been observed.⁹⁵

Treatment and management

Currently, there are no specific anti-MERS-CoV therapies available for human use. The mainstay of clinical management of MERS cases is mostly symptom focused, providing supportive care^{10,61} with pain and fever management, treating concomitant or secondary bacterial infections, and supporting vital organ functions. Mild MERS cases can be managed at home.⁹⁶ Predictive factors for development of pneumonia include older age (>56 years), high fever, thrombocytopenia, lymphopenia, C-reactive protein greater than or equal to 2 mg/dL, and a high viral load in sputum (threshold cycle value of rRT-PCR <28.5).^{10,70} Respiratory failure and acute kidney injury (with haematuria and proteinuria) are common among patients admitted for hospital care because of the severity of their disease and who require mechanical ventilation, extracorporeal membrane oxygenation, and dialyses.^{10,61,70}

Several empirical treatments have been studied in severely ill patients. Therapies used for severely ill patients with MERS have included convalescent plasma, corticosteroids, antiviral agents (eg, interferons and ribavirin), and protease inhibitors, or combination of members from these groups. A systematic review of 30 publications on therapeutic agents used for MERS treatment in various outbreaks showed a complete absence of an accurate evidence base and emphasised the need for controlled trials.⁹⁷

Antiviral agents

Several agents have shown inhibitory effects against MERS-CoV in cell cultures, including interferons, ribavirin, ciclosporin, and mycophenolic acid.^{98–105} Empirical lopinavir–ritonavir, pegylated interferon alfa-2a, and

ribavirin have been used for serious cases, but no efficacy data are yet available. There is an ongoing randomised clinical trial in progress in Saudi Arabia (the MIRACLE trial; NCT02845843) comparing lopinavir–ritonavir, recombinant interferon beta, and standard supportive care against placebo and standard supportive care in patients with laboratory-confirmed MERS requiring hospital admission. This recursive, multicentre, placebo-controlled, double-blind, randomised controlled trial is designed to include two components, each consisting of two stages. The first two-stage component is designed to adjust sample size and determine futility stopping, but not efficacy stopping (n=34). The second two-stage component is designed to determine efficacy stopping and possibly readjustment of sample size.

Convalescent plasma

The use of plasma from patients with MERS who have made a full recovery (convalescent plasma) requires clinical trial evaluation. Preclinical animal data show that transfer of serum samples from MERS-CoV-immune camels to infected mice resulted in reduced weight loss and lung histopathology,¹⁰⁶ thus suggesting therapeutic potential. In the 2015 Korean outbreak, three of 13 patients with MERS with respiratory failure were given four infusions of convalescent plasma.¹⁰⁷ Two of three showed neutralising activity. Donor plasma with a plaque reduction neutralisation test (PRNT) titre of 1/80 had a meaningful serological response after convalescent plasma infusion, while that with a PRNT titre of 1/40 did not. The authors conclude that for effective convalescent plasma infusion in patients with MERS, donor plasma with a neutralisation activity of a PRNT titre greater than or equal to 1/80 should be used.

Antibiotic therapy

Antibiotic therapy is commonly started empirically in seriously ill patients. A retrospective study of 136 patients with MERS found that macrolide therapy resulted in no reduction in mortality or faster MERS-CoV RNA clearance compared with those who were not treated with macrolides.¹⁰⁸

Intensive care management

For serious cases, hospital inpatient care is required to reduce the risk of complications such as organ failure and secondary infections. Non-invasive ventilation is associated with a high failure rate (92%) in patients with acute hypoxaemic respiratory failure due to MERS-CoV infection.¹⁰⁹ Patients with severe symptoms might need to be managed in an intensive care unit, where lung-protective ventilatory strategies for acute respiratory distress syndrome, inotropic support, antimicrobial therapy for co-infections, and renal replacement therapy for acute renal failure can be provided.

Panel 3: Potential therapies for the treatment of MERS coronavirus infection^{97-105, 139-143}**Antibody-based interventions**

- Whole blood
- Convalescent plasma
- Intravenous immunoglobulin
- Polyclonal human immunoglobulin (SAB-301) from transgenic cows
- Equine antibody fragments
- Camel antibodies
- Monoclonal antibodies (eg, MERS-4, MERS-7)
- Human monoclonal antibodies to Sprotein
- Humanised anti-S monoclonals (eg, hMS-1, m336, 4C2)

Interferons

- Interferon alfa (1a, 2b)
- Interferon beta-1b
- Interferon gamma

Antivirals

- Ribavirin monotherapy (with or without interferon)
- HIV protease inhibitors (eg, lopinavir, nelfinavir)
- Cyclophilin inhibitors (eg, ciclosporin, alisporivir)
- Nucleoside viral RNA polymerase inhibitors (galidesivir, remdesivir, EIDD 2801 [prodrug of EIDD 1931])

MERS-CoV 3C-like protease inhibitors

- Flavonoids (eg, herbacetin, isobavachalcone, quercetin 3-β-d-glucoside, helichrysetin)

Combination therapies

- Lopinavir-ritonavir and interferon beta 1b
- Cyclosporin plus interferon alpha
- Lopinavir-ritonavir
- Ribavirin and interferon alpha

Host-directed therapies (repurposed drugs)

- Chlorpromazine
- Mycophenolic acid (with or without interferon beta 1b)
- Nitazoxanide
- Sitagliptin
- Omacetaxine mepesuccinate
- Aciclovir
- Imatinib mesylate (tyrosine mesylate inhibitor)
- Neurotransmitter inhibitors (eg, clomipramine, astemizole)
- Neutriceuticals (eg, zinc)

Host-directed therapies (cellular therapy)

- Allogeneic mesenchymal stromal cells

Lectins

- Mannose binding lectin

Anti-coronavirus peptides

- Peptides derived from HR1, HR2, and RBD spike protein subunits
- Peptides inhibiting viral entry and replication
- Peptides derived from antimicrobial peptides

RNA interference molecules silencing key MERS-CoV genes

- miRNA molecules
- siRNA molecules

MERS=Middle East respiratory syndrome. MERS-CoV=Middle East respiratory syndrome coronavirus. RBD=receptor-binding domain. S=spike. RNA=ribonucleic acid. miRNA=micro RNA. siRNA=small interfering RNA.

Systemic corticosteroids

Studies show no benefit from the use of systemic corticosteroids. Systemic corticosteroids were shown to delay viral clearance in critically ill patients with MERS-CoV infection.¹⁰⁰

New treatments

A range of potential treatments are under consideration, development, or evaluation (panel 3). A phase 1, first-in-human, single site, randomised, double-blind, placebo-controlled study to evaluate the safety, tolerability, pharmacokinetics, and immunogenicity of single ascending doses of a co-administered monoclonal antibodies REGN3048 and REGN3051 administered intravenously in healthy adult volunteers (NCT03301090) is ongoing and study duration will be approximately 16 months. Another study is comparing patients who received extracorporeal membrane oxygenation (ECMO) support with those who did not receive ECMO support (NCT02627378). Several monoclonal and polyclonal neutralising antibody products, including novel chimeric camel or human heavy-chain antibodies, are protective in animal models,¹¹⁰⁻¹¹² and their potential use for pre-exposure or post-exposure prophylaxis in outbreak management requires evaluations in controlled trials.

Mental health support

In a mental health study, 7·6% of 1656 patients who were quarantined showed symptoms of anxiety, and 6·4% reported feelings of anger during the 2 weeks of quarantine.¹¹³ Mental health support, accurate information, and appropriate supplies, including food, clothes, and accommodation, should be provided to people who feel isolated.

Management outcomes and mortality rates

Several studies have reported disease severity and mortality risk factor data from patients in the Middle East and South Korea.^{11,12,21,114-117} Ahmed and colleagues collected the daily information on MERS-CoV cases posted online by the Saudi Arabian Ministry of Health between Dec 2, 2014, and Nov 12, 2016, and reviewed 660 laboratory-confirmed cases of MERS.¹¹⁴ They showed mortality at day 3 (13·8%), day 30 (28·3%), and overall (29·8%). Patients older than 60 years were more likely to die (45·2% mortality) from their infections than were younger patients (20%).

Several factors are associated with severe disease and high mortality rates in patients with MERS. These factors include male sex, comorbid pre-existing illnesses (such as obesity, diabetes mellitus, cancer, chronic heart, lung, and kidney disease, and immunocompromised states), low serum albumin, thrombocytopenia, lymphopenia, concomitant infections, and positive plasma MERS-CoV RNA. DPP4 receptors are upregulated in the lungs of smokers and patients with chronic obstructive pulmonary disease, which might explain why patients with comorbid

Panel 4: Priorities for MERS research, surveillance, management, and control

Establishing a One Health global consortium:

- Creation of a One Health multidisciplinary consortia in Middle Eastern and African countries to tackle zoonotic spread of MERS-CoV between animals, humans, and the environment and to monitor, respond to, and prevent major outbreaks
- Establishing effective One Health research and surveillance collaborations between a range of expertise including public health officers, physicians, veterinarians, animal husbandry specialists, agriculturalists, ecologists, phylogeneticists, trialists, and basic scientists

Priority studies in dromedary camels:

- Natural history studies that evaluate evidence of re-infection
- Value chain and production system analyses
- Improving surveillance to evaluate any seasonal or temporal variation in viral shedding
- Identifying important points for interventions and interruption of within-species and zoonotic transmission
- Accelerating the development of camel vaccine candidates

Priority studies at the animal-human-environmental interface:

- Real-time mapping of virus circulation and geographical range of MERS-CoV infection in humans and dromedary camels
- Development of new accurate and accessible point-of-care diagnostic tests for outbreak containment, case management, and surveillance
- Evaluating geographical extent of spillover from animals to humans in Africa, the Middle East, and south Asia
- Animal and human serological, virological, and genomic studies in specific locations to evaluate risk factors for human infection and exact routes of zoonotic transmission, including food and oral routes
- Social science and anthropological studies to describe and quantify exposures to dromedary camels and identify opportunities for risk-mitigating interventions

Priority studies in human populations:

- Accelerating research and development, implementation, and evaluation of medical countermeasures to reduce morbidity and mortality associated with MERS (especially in high risk comorbid populations)
- Identifying optimal interventions for reducing infection among health-care workers in hospital settings
- Determining the role of asymptomatic cases in transmission of infections in humans and whether any specific behaviours can result in human infection from non-human sources
- Targeted epidemiological studies in clinical settings to better understand immune response and duration of infectiousness and address the long-term medical and psychological complications in MERS survivors
- Integrating testing for MERS-CoV into existing respiratory disease surveillance systems to identify extent and spectrum of mild infection in the community

Priority studies for vaccines, therapeutics, and rapid diagnostics:

- Accelerating the development of human MERS-CoV vaccine candidates
- Accelerating development of MERS-CoV therapeutic pipeline and host-directed therapies
- Development of new accurate and accessible diagnostic tests for outbreak containment, case management, and surveillance
- Increased donor investment into research and development for: novel point-of-care testing options, syndromic panels for differential diagnosis, a greater understanding of viral and antibody kinetics, improved access to clinical specimens, and establishment of international reference standards

Adapted from the FAO-OIE-WHO MERS Technical Working Group's report on MERS (2018). MERS=Middle East respiratory syndrome. ³⁵ MERS-CoV=Middle East respiratory syndrome coronavirus.

lung diseases are prone to severe illness.¹¹⁸ In the 2015 Korean outbreak, the case-fatality rate was 20·4% (39 of 186 patients died), the in-hospital mortality was 19·4% (36 of 186 patients), 7-day mortality (from symptom onset) was 3·8% (seven of 186), and 28-day mortality was 17·7% (33 of 186).^{11,12} Host factors associated with mortality in this outbreak were older age (>60 years), smoking history, pre-existing pneumonia, abnormal renal function, and comorbidity. Low albumin, altered mentality, and high pneumonia severity index score at admission were risk factors for mortality.

Prognosis and long term follow-up of MERS survivors

In a study of long-term effects after recovery from MERS,¹¹⁹ patients with severe illness who were admitted to intensive care reported higher limitations in some

measure of their quality of life than patients with less severe illness who were managed in medical wards. The long-term consequences of MERS illness on survivors appear to be similar to those caused by other viral causes of severe acute respiratory infections such as influenza, parainfluenza, and metapneumovirus.¹²⁰ Follow-up imaging studies of patients who recover from MERS indicate that lung fibrosis might develop in a substantial number of patients.⁹⁵

Prevention

The prevention of transmission of MERS-CoV in the community and in health-care settings is crucial to preventing outbreaks and further spread. Several substantive reviews and WHO guidelines are available on the subject.^{21,61,76,96}

It is important to maintain good personal and environmental hygiene, and to implement stringent contact and droplet precautions among health-care workers. To prevent community transmission, contact tracing, quarantine or isolation of close contacts, and public education are important measures.^{20,60}

In hospitals, early case detection followed by isolation is essential, ideally in negative pressure isolation rooms. The main infection prevention and control measures for managing patients with MERS are well documented from the SARS epidemic and from experiences from managing MERS outbreaks (appendix pp 7–8).

Advances in vaccine development

Advances in technology, vaccine platforms, clinical trial designs, and bioinformatics are supporting MERS-CoV vaccine development (appendix pp 8–9). WHO target product profiles for MERS-CoV vaccines calls for the development of three types of MERS vaccines: a human vaccine for long-term protection of people at high exposure risk, such as health-care workers and those working with potentially infected camels; a human vaccine for reactive use in outbreak settings; and a dromedary camel vaccine to prevent zoonotic transmission.¹²¹ Multiple types of vaccine candidates are in development including inactivated whole virus, live attenuated virus, viral vectored vaccines, subunit vaccines, and DNA vaccines.^{121,122} Most vaccines use the S protein or the domain of the S protein required for binding to host DPP4 as an immunogen, since neutralising antibodies are mostly directed to the receptor binding domain.

Priorities for MERS research, surveillance, management, and control

WHO, Food and Agriculture Organization of the UN, and World Organization for Animal Health, in consultation with global MERS community, identified knowledge gaps and priorities for MERS research, surveillance, management, and control (panel 4).³¹

Several recent publications have highlighted the importance of a one-human-environmental-animal-health (One Health) approach to tackle and control the spread of MERS-CoV (appendix p 9).^{31,121–126}

Conclusion

MERS-CoV is a pathogen with epidemic potential that continues to cause sporadic human disease and remains on the WHO Blueprint 2020 priority list.⁴ Given that MERS-CoV appears to be highly endemic among dromedary camels from geographically widespread areas of the Middle East and Africa, zoonotic transmission with consequent risk of human epidemics will most likely continue for years to come.^{8,9} MERS-CoV endemic and at-risk countries must invest more in surveillance, in public health research, and in medical interventions—including human and camel vaccine development. The continued risk of human MERS-CoV outbreaks 7 years

after its first discovery, effective human and camel MERS-CoV vaccines appear to be the ideal way to prevent continuing spread of MERS-CoV in dromedary camels in the Middle East and in humans at high risk of acquiring community and nosocomial MERS-CoV infection.

Contributors

This paper was conceptualised by ZAM and AZ. All authors contributed equally to the literature search, writing, and reviewing of the manuscript.

Declaration of interests

We declare no competing interests.

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