

# Middle East Respiratory Syndrome (MERS) coronavirus seroprevalence in domestic livestock in Saudi Arabia, 2010 to 2013

M G Hemida<sup>1,2,3</sup>, R A Perera<sup>1,4,5</sup>, P Wang<sup>6,7</sup>, M A Alhamadi<sup>2</sup>, L Y Siu<sup>5,6</sup>, M Li<sup>5,6</sup>, L L Poon<sup>4,5</sup>, L Saif<sup>8</sup>, A Alnaeem (aalnaeem@kfu.edu.sa)<sup>9</sup>, M Peiris (malik@hku.hk)<sup>4,5,6</sup>

1. These authors contributed equally to this work

2. Department of Microbiology and Parasitology, College of Veterinary Medicine, King Faisal University, Saudi Arabia

3. Department of Virology, Faculty of Veterinary Medicine, Kaferelshiek University, Egypt

4. Centre of Influenza Research, The University of Hong Kong, Hong Kong SAR, China

5. School of Public Health, The University of Hong Kong, Hong Kong SAR, China

6. HKU-Pasteur Pole, The University of Hong Kong, Hong Kong SAR, China

7. Key Laboratory of Protein and Peptide Pharmaceuticals, Chinese Academy of Sciences - University of Tokyo Joint Laboratory of Structural Virology and Immunology, Beijing, China

8. Food Animal Health Research Program, Veterinary Preventive Medicine Department, OARDC/The Ohio State University, Wooster, United States of America

9. Department of Clinical Studies, College of Veterinary Medicine, King Faisal University, Saudi Arabia

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**In Saudi Arabia, including regions of Riyadh and Al Ahsa, pseudoparticle neutralisation (ppNT) and microneutralisation (MNT) tests detected no antibodies to Middle East Respiratory Syndrome coronavirus (MERS-CoV) in sheep (n= 100), goats (n= 45), cattle (n= 50) and chickens (n= 240). Dromedary camels however, had a high prevalence of MERS-CoV antibodies. Bovine coronavirus (BCoV) infected sera from cattle had no cross-reactivity in MERS-CoV ppNT or MNT, while many dromedary camels' sera reacted to both BCoV and MERS-CoV. Some nevertheless displayed specific serologic reaction profiles to MERS-CoV.**

In a seroepidemiological study of domestic livestock (sheep, goats, cattle, chicken) and dromedary camels from Saudi Arabia, we find that only dromedary camels have evidence of seropositivity to Middle East Respiratory Syndrome coronavirus (MERS-CoV), suggesting an infection with a MERS-CoV-like virus. Although some dromedary camels examined had specific serologic reaction profiles to MERS-CoV, many had sera displaying reactivity to both MERS-CoV and bovine coronavirus (BCoV) and the sera appear to have an unusually broad pattern of cross-reactivity for these related viruses. This needs to be considered when interpreting seroepidemiological data by carrying out parallel microneutralisation tests for both MERS CoV and BCoV.

## Middle East Respiratory Syndrome coronavirus

MERS coronavirus was recognised as a cause of severe human respiratory disease in 2012 [1]. As of 22 November 2013, 157 laboratory-confirmed, and 19

other probable cases of MERS have been reported, 69 of these being fatal [2]. Sporadic or index cases account for approximately 40% of the cases and have occurred in Jordan, Kuwait, Oman, Qatar, Saudi Arabia and United Arab Emirates. Imported cases have been reported in France, Germany, Italy, Spain, Tunisia and United Kingdom, with secondary transmission documented in some of these cases [2,3]. Human-to-human transmission has occasionally been observed, the largest cluster of such secondary transmission being reported from Al Ahsa, Saudi Arabia [4]. Phylogenetic analysis suggests considerable diversity within MERS-CoV sequences analysed to date, with two distinct clades of virus being noted [5].

Index cases of MERS appear to be zoonotic in origin, with contact with domestic livestock (e.g. camels, sheep) being reported in some of these patients [3]. But neither the proximate animal source of human infection nor the natural reservoir of the virus is known. Closely related, but not identical viruses have been found in species of insectivorous bats [6]. There is a preliminary report of a short fragment of virus genome almost identical to MERS-CoV reportedly found in Egyptian tomb bats [7].

Two recent studies from Oman and Egypt respectively, two Middle-Eastern countries adjacent to affected ones, reported high rates of MERS-CoV seroprevalence in dromedary camels [8,9]. However, there have been no seroepidemiological data of domestic livestock, camels or wildlife from affected countries, to date.

## FIGURE

Locations where serum samples were collected for the Middle East Respiratory Syndrome (MERS) coronavirus seroepidemiological study in domestic livestock, Saudi Arabia, 2010–2013



### Seroepidemiological study

Serum samples were collected from dromedary camels (n= 310), sheep (n= 100), goats (n= 45), cattle (n= 50) and chicken (n= 240) from Riyadh, Al Ahsa and other regions of Saudi Arabia (Figure).

The MERS-CoV serology methods used have been described in a previous publication [9]. The sera were heat inactivated at 56 °C for 30 minutes and screened for antibody to MERS-CoV at a serum dilution of 1:20 in a MERS-CoV pseudoparticle neutralisation test (ppNT) [9]. Camel sera (n=56) that were seropositive at a screening dilution of 1:20 in the ppNT assay were randomly selected, representing each age group, less than one year (n=15), one to three years (n=12), four to five years (n=16) and older than five years (n=13), and

titrated to end point in the MERS-CoV ppNT, MERS-CoV microneutralisation test (MNT) and bovine coronavirus (BCoV) MNT.

The MNTs were done using MERS-CoV (strain: EMC) obtained from Dr R Fouchier, Erasmus MC, Rotterdam and bovine coronavirus (ATCC BRCV-OK-0514-2). The MNTs were done on Vero cells (ATCC CCL-81) for MERS-CoV and HRT-18G cells (obtained from ATCC) for BCoV, respectively. Serial two-fold dilutions of heat-inactivated sera (56 °C for 30 minutes) were mixed with equal volumes of 200 tissue culture infective dose (TCID)<sub>50</sub> of virus and incubated for one hour at 37 °C. The virus–serum mixture was then added in quadruplicate to cell monolayers in 96-well microtitre plates. After one hour of adsorption, the virus-serum mixture

**TABLE 1**

Location and time of sampling of animals chosen for MERS-CoV ppNT and proportion of serum samples positive at 1:20 dilution, Saudi Arabia, 2010–2013 (n=745 samples)

Animals	Location (number of animals)	Collection period	Serum samples collected N	Sera positive in MERS-CoV ppNT N (%)
Dromedary camels	Al-Ahsa (120), Riyadh (190)	2012–2013	310	280 (90)
Sheep	Al-Ahsa (100)	2012–2013	100	0 (0)
Goats	Al-Ahsa (15), Taif (10), Madinah (10), Qatif (10)	2010–2012	45	0 (0)
Chicken	Al-Ahsa (120), Dammam and Alkhober (80), Abqaiq (40)	2012–2013	240	0 (0)
Cattle	Al-Ahsa (17), Taif (13), Madinah (10), Qatif (10)	2010–2013	50	0 (0)

MERS-CoV ppNT: Middle East Respiratory Syndrome pseudoparticle neutralisation test.

Riyadh province is in central Saudi Arabia; Al Ahsa, Qatif, Dammam, Alkhober and Abqaiq are in the eastern part of Saudi Arabia; Madinah and Taif are in the western part of Saudi Arabia (Figure).

was removed and 150µl of fresh culture medium was added to each well and the plates incubated at 37 °C in 5% CO<sub>2</sub> in a humidified incubator. A virus back-titration was performed without immune serum to assess input virus dose.

Cytopathic effect (CPE) was read at three days post infection for MERS-CoV and four days post infection for BCoV. The highest serum dilution that completely protected the cells from CPE in half of the wells was defined as the neutralising antibody titre. Positive and negative control sera were included in each assay.

As positive controls, we used a convalescent serum from a human patient with MERS kindly provided by Dr C Drosten, Institute of Virology, University of Bonn Medical Centre, Bonn (MERS-CoV ppNT 1:160); and sera from two experimentally infected macaques (ppNT

titres 1:40) and a non-infected control macaque (ppNT <1:20) kindly provided by Bart Haagmans, Erasmus MC as reported previously [9]. Additional positive and negative controls used were sera from dromedary camels from Egypt previously found to be seropositive (ppNT 1:320) and sero-negative (ppNT<1:20) [9]. Antisera to BCoV from an experimentally inoculated gnotobiotic calf and guinea pig were provided by Dr. Linda Saif.

## Results

Sera that were positive in the MERS-CoV ppNT test at a screening dilution of 1:20 are shown in Table 1.

A gnotobiotic calf and guinea pig B CoV antisera that had a homologous reaction titre of ≥1:1,280 and 1:80 respectively, did not cross-neutralise MERS-CoV in either the MERS-CoV ppNT or MNT. None of the sheep, goat, cattle or chicken sera had any MERS-CoV ppNT

**TABLE 2**

Stratification by age and location of dromedary camels testing positive in the MERS-CoV ppNT assay at a serum dilution of 1:20, Saudi Arabia, 2012–2013 (n=310 animals)

Age group (years)	Riyadh		Al Ahsa		Overall	
	Animals tested N	Animals testing positive N (%)	Animals tested N	Animals testing positive N (%)	Animals tested N	Animals testing positive N (%)
<1	21	18 (86)	44	29 (66)	65	47 (72)
1–3	55	52 (94)	51	49 (96)	106	101 (95)
4–5	52	51 (98)	24	23 (96)	76	74 (97)
>5	62	57 (92)	1	1 (100)	63	58 (92)
Total	190	178 (94)	120	102 (85)	310	280 (90)

MERS-CoV ppNT: Middle East Respiratory Syndrome pseudoparticle neutralisation test.

TABLE 3A

Serology titres for MERS-CoV and BCoV in dromedary camel sera, Saudi Arabia, 2012–2013 (n=56)

Sera IDs	Age group in years	MERS-CoV ppNT titre	MERS-CoV MNT titre	BCoV MNT titre	Reaction profile	Region
1	<1	1:2,560	1:160	1:20	MERS-CoV specific	Riyadh
2	<1	1:320	1:20	<1:10	MERS-CoV specific	Riyadh
3	<1	≥1:5,120	1:640	1:40	MERS-CoV specific	Al-Ahsa
4	<1	1:1,280	1:320	<1:10	MERS-CoV specific	Al-Ahsa
5	<1	1:640	1:40	1:160	BCoV specific	Riyadh
6	<1	1:640	1:40	1:320	BCoV specific	Riyadh
7	<1	1:160	1:10	1:40	BCoV specific	Riyadh
8	<1	1:1,280	1:80	1:640	BCoV specific	Al-Ahsa
9	<1	<1:20	<1:10	1:160	BCoV specific	Al-Ahsa
10	<1	1:80	1:10	1:40	BCoV specific	Al-Ahsa
11	<1	1:80	1:10	<1:10	Indeterminate	Riyadh
12	<1	≥1:5,120	1:640	1:640	Indeterminate	Riyadh
13	<1	1:2,560	1:320	1:320	Indeterminate	Al-Ahsa
14	<1	1:2,560	1:320	1:320	Indeterminate	Al-Ahsa
15	<1	1:1,280	1:160	1:320	Indeterminate	Al-Ahsa
16	1–3	≥1:5,120	≥1:1,280	<1:10	MERS-CoV specific	Riyadh
17	1–3	1:160	1:20	<1:10	MERS-CoV specific	Riyadh
18	1–3	≥1:5,120	≥1:1,280	1:80	MERS-CoV specific	Riyadh
19	1–3	1:640	1:80	<1:10	MERS-CoV specific	Al-Ahsa
20	1–3	1:640	1:160	<1:10	MERS-CoV specific	Al-Ahsa
21	1–3	1:640	1:40	1:160	BCoV specific	Riyadh
22	1–3	1:20	<1:10	1:160	BCoV specific	Riyadh
23	1–3	1:2,560	1:80	≥1:1,280	BCoV specific	Al-Ahsa
24	1–3	1:320	1:10	1:20	Indeterminate	Riyadh
25	1–3	1:640	1:80	1:40	Indeterminate	Riyadh
26	1–3	1:160	1:20	1:40	Indeterminate	Al-Ahsa
27	1–3	1:1,280	1:640	1:320	Indeterminate	Al-Ahsa
28	4–5	≥1:5,120	1:640	1:160	MERS-CoV specific	Riyadh
29	4–5	1:160	1:160	1:40	MERS-CoV specific	Riyadh
30	4–5	≥1:5,120	1:320	1:80	MERS-CoV specific	Riyadh
31	4–5	≥1:5,120	1:640	1:80	MERS-CoV specific	Riyadh
32	4–5	≥1:5,120	≥1:1,280	1:40	MERS-CoV specific	Riyadh
33	4–5	1:640	1:160	<1:10	MERS-CoV specific	Riyadh
34	4–5	≥1:5,120	≥1:1,280	<1:10	MERS-CoV specific	Al-Ahsa
35	4–5	1:320	1:40	1:640	BCoV specific	Riyadh
36	4–5	1:1,280	1:320	≥1:1,280	BCoV specific	Riyadh
37	4–5	1:2,560	1:640	≥1:1,280	BCoV specific	Riyadh
38	4–5	1:320	1:40	1:320	BCoV specific	Al-Ahsa
39	4–5	≥1:5,120	1:320	≥1:1,280	BCoV specific	Al-Ahsa
40	4–5	1:160	1:40	1:320	BCoV specific	Al-Ahsa
41	4–5	1:1,280	1:160	1:320	Indeterminate	Riyadh
42	4–5	1:2,560	1:320	1:640	Indeterminate	Riyadh
43	4–5	1:1,280	1:160	1:320	Indeterminate	Al-Ahsa

BCoV: bovine coronavirus; ID: identity; MERS-CoV: Middle East Respiratory Syndrome coronavirus; MNT: microneutralisation test; ppNT: pseudoparticle neutralisation test.

The reaction profile was denoted as MERS-CoV specific if the antibody titre to MERS-CoV was ≥4 fold higher than BCoV; and BCoV specific if the titre to BCoV was ≥4 fold higher than for MERS-CoV. Other sera are regarded to have an indeterminate reaction profile.

**TABLE 3B**

Serology titres for MERS-CoV and BCoV in dromedary camel sera, Saudi Arabia, 2012–2013 (n=56)

Sera IDs	Age group in years	MERS-CoV ppNT titre	MERS-CoV MNT titre	BCoV MNT titre	Reaction profile	Region
44	>5	≥1:5,120	1:640	1:80	MERS-CoV specific	Riyadh
45	>5	1:1,280	1:320	1:80	MERS-CoV specific	Riyadh
46	>5	1:640	1:640	1:40	MERS-CoV specific	Riyadh
47	>5	<1:20	<1:10	1:80	BCoV specific	Riyadh
48	>5	1:80	1:80	1:320	BCoV specific	Riyadh
49	>5	≥1:5,120	1:640	>1:1280	Indeterminate	Riyadh
50	>5	1:1,280	1:320	1:160	Indeterminate	Riyadh
51	>5	1:1,280	1:640	1:320	Indeterminate	Riyadh
52	>5	≥1:5,120	1:640	>1:1,280	Indeterminate	Riyadh
53	>5	1:1,280	1:640	1:320	Indeterminate	Riyadh
54	>5	1:640	1:640	1:640	Indeterminate	Riyadh
55	>5	≥1:5,120	≥1:1,280	>1:1,280	Indeterminate	Riyadh
56	>5	1:320	1:20	1:40	Indeterminate	Riyadh

BCoV: bovine coronavirus; ID: identity; MERS-CoV: Middle East Respiratory Syndrome coronavirus; MNT: microneutralisation test; ppNT: pseudoparticle neutralisation test.

The reaction profile was denoted as MERS-CoV specific if the antibody titre to MERS-CoV was ≥4 fold higher than BCoV; and BCoV specific if the titre to BCoV was ≥4 fold higher than for MERS-CoV. Other sera are regarded to have an indeterminate reaction profile.

activity while 280 of 310 (90%) dromedary camel sera were seropositive. The age-group and location of the camels tested and their serostatus in the screening assay are shown in Table 2. While 47 of 65 (72%) of camels less than one year of age were seropositive, 233 of 245 (95%) of camels older than one year were seropositive to MERS-CoV in the ppNT test (chi-squared test,  $p < 0.01$ ).

A randomly selected subset of 54 ppNT seropositive sera and two ppNT negative sera were titrated by MERS-CoV ppNT, MERS-CoV MNT and BCoV MNT (Table 3). MERS-CoV ppNT titres ranged from 1:20 to ≥1:5,120 and MERS-CoV MNT titres ranged from <1:10 (in one serum only) to ≥1:1,280. Dromedary camel sera, which have ≥4 fold higher antibody MERS-CoV MNT titres compared to BCoV MNT titres are denoted as ‘MERS-CoV specific’ reaction patterns. Sera with comparable (within 4-fold) titres to both viruses and are denoted as ‘indeterminate’ reaction patterns while those that have ≥4 fold higher titres to BCoV are regarded as ‘BCoV specific’.

Twelve of the 50 cattle sera were randomly selected for BCoV MNT and 11 sera were positive (Table 4).

## Discussion

This study was conducted Saudi Arabia, including in central and eastern provinces from which most of the human cases of MERS-CoV hitherto had been

detected; in particular, the Al Ahsa region from where the largest cluster of human-to-human transmission of the disease has occurred [4] and the Riyadh region, which appears to host the greatest genetic diversity of MERS-CoV within Saudi Arabia [5]. None of the cattle, goat, sheep or chickens had any detectable antibody to MERS-CoV. While larger numbers of such livestock in the vicinity of confirmed cases of MERS need to be examined, our data do not indicate that these species of domestic livestock are commonly infected by MERS-CoV. As expected, BCoV antibody was common in cattle with 11 of 12 cattle sera having evidence of BCoV MNT antibody, but these sera did not cross-react with MERS-CoV.

Dromedary camels from the central province of Riyadh as well as the eastern province, Al Ahsa, had comparably high levels of seropositivity to MERS-CoV by ppNT tests. Camels less than one year of age had lower seroprevalence (72%) than those older than one year of age (95%) ( $p < 0.01$ ). Some dromedary camels have high titres to MERS-CoV in the absence of any reactivity to BCoV confirming that these animals are being infected by a virus very different to BCoV, which could be identical or closely related to MERS-CoV [8,9]. Similarly, some animals appear to be infected by a BCoV-like virus without cross-reactivity to MERS-CoV. If we consider just the ‘MERS-CoV specific’ reactions, we observe that these animals are getting infected within the first year of life (Table 3).

**TABLE 4**

Serology titres for MERS-CoV and BCoV in cattle sera, Saudi Arabia, 2010–2013 (n=12)

Sera ID	MERS-CoV ppNT titre	MERS-CoV MNT titre	BCoV MNT titre
1	<1:20	<1:10	1:40
2	<1:20	<1:10	<1:10
3	<1:20	<1:10	1:640
4	<1:20	<1:10	1:160
5	<1:20	<1:10	1:80
6	<1:20	<1:10	1:40
7	<1:20	<1:10	1:160
8	<1:20	<1:10	1:40
9	<1:20	<1:10	1:40
10	<1:20	<1:10	1:640
11	<1:20	<1:10	1:160
12	<1:20	<1:10	1:160

BCoV: bovine coronavirus; ID: identity; MERS-CoV: Middle East Respiratory Syndrome coronavirus; MNT: microneutralisation test; ppNT: pseudoparticle neutralisation test.

Naturally infected cattle with BCoV titres of up to 1:640 and an experimentally infected calf with a titre of  $\geq 1:1,280$  had no cross-reaction to MERS-CoV in ppNT and MNT assays. Previously, we had shown that human sera with antibody to Severe Acute Respiratory Syndrome (SARS)-CoV failed to react with MERS-CoV, moreover SARS-CoV was not neutralised by dromedary camel sera with high antibody titres to MERS-CoV [9]. In contrast, while some dromedary camel sera had mono-specific reactions to either MERS CoV or BCoV, many sera have ‘indeterminate’ reaction profiles with reactivity to both MERS-CoV and BCoV (Table 3). It is possible that sequential infection of dromedary camels with different coronaviruses may lead to a broadening of the serological cross-reaction profile. Alternatively, it is possible that the unusual single chain immunoglobulins possessed by camels [10] may give broader serological cross-reactivity than is seen with other species. In any event, these findings highlight the importance of parallel titration of sera to both MERS-CoV and BCoV when interpreting seroepidemiological results.

The exact identity of this MERS-CoV-like virus can only be established by characterising this virus from dromedary camel specimens. As many of the dromedary camels sampled in this study were already seropositive to MERS-CoV, we did not attempt to detect viral RNA in the serum samples. There are (as yet unpublished) reports of detecting a MERS-CoV-like virus in camel specimens from Saudi Arabia and Qatar [2]. Assuming that MERS-CoV seropositivity in MNTs, irrespective of the infecting virus that leads to such seropositivity, correlates with

resistance to, or reduced susceptibility to re-infection, these results would suggest that the search for virus in these animals should be focused on dromedary camels <1 year of age. Studies involving follow-up of herds of camels from time of calving through the first year of life with serial blood samples together with oral and rectal or fresh faecal swabs would better define the ecology of the MERS-CoV-like virus infecting these animals and provide virus for genetic characterisation. Such studies are a priority to determine whether dromedary camels are in fact a source of human MERS-CoV infection or whether they are being infected by a ubiquitous novel coronavirus closely related to MERS-CoV. As illustrated with SARS a decade ago and avian influenza A(H7N9) recently, identification of the animal source and the setting within which zoonotic transmission occurs can provide options for reducing repeated zoonotic transmissions and enhancing global public health. Finally, given the high titres of neutralising antibody to MERS CoV commonly seen in dromedary camels, serum from these animals may provide an option for passive immunotherapy of patients with MERS, from whom no specific antiviral therapy currently exists.

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#### Conflict of interest

None declared.

#### Authors' contributions

Maged Gomaa Hemida (mhemida@kfu.edu.sa), Malik Peiris (malik@hku.hk) designed, coordinated and supervised the study and wrote the manuscript. Ranawaka APM Perera (mahenp@hku.hk) developed the MERS-CoV and bovine coronavirus microneutralisation tests, carried out the MERS-CoV tests in BSL3 containment and the MERS pseudoparticle assays. Pei-gang Wang (pgwang@hkucc.hku.hk) developed the MERS-CoV pseudotype assay. Mohamed A Alhammadi (malhammadi@kfu.edu.sa): Specimen collection and data recording. Lewis YL Siu (ylsiu@hku.hk) and Ming-yuan Li (lmy288@hku.hk) carried out the MERS-CoV pseudoparticle assays. Leo LLM Poon (llmpoon@hku.hk): Provided advice on laboratory methods. Linda Saif (saif.2@osu.edu): Provided reagents and advised on study design. Abdelmohsen Alnaeem (aalnaeem@kfu.edu.sa): Designed and supervised the study. All authors reviewed and commented on the manuscript.

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