

THE LANCET

Supplementary appendix

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Supplement to: Guery B, Poissy J, el Mansouf L, et al, and the MERS-CoV study group et al. Clinical features and viral diagnosis of two cases of infection with Middle East Respiratory Syndrome coronavirus: a report of nosocomial transmission. *Lancet* 2013; published online May 29. [http://dx.doi.org/10.1016/S0140-6736\(13\)60982-4](http://dx.doi.org/10.1016/S0140-6736(13)60982-4).

Figure 1S

RdRp

	20	40	60	80					
EMC	TCGCACTGTT	GCAGGGCGTGT	CCATACTTAG	CACAATGACT	AATGCCAGT	ACCATCAGAA	AATGCTTAAG	TCCATGGCTG	80
JOR	80
ENG1	80
ENG/QA	80
MUN/ABD	80
FR1/UAE	80
FR2	80
	100	120	140	160					
EMC	CAACTCGTGG	AGCGACTTGC	GTCATTGGTA	CTACAAAGTT	CTACGGTGGC	TGGGATTTCA	TGCTTAAAAC	ATTGTACAAA	160
JOR	T	.	T	160
ENG1	t	.	T	160
ENG/QA	T	.	T	160
MUN/ABD	T	.	T	160
FR1/UAE	T	.	T	160
FR2	T	.	T	160
	180								
EMC	GATGTTGATA	ATCCGCATCT	TA	182					
JOR	.	.	.	182					
ENG1	.	.	.	182					
ENG/QA	.	.	.	182					
MUN/ABD	.	.	.	182					
FR1/UAE	.	.	.	182					
FR2	.	.	.	182					

N

	20	40	60	80					
EMC	AATAAGTGGT	TGGAGCTTCT	TGAGCAAAT	ATTGATGCCT	ACAAAAACCTT	CCCTAAGAAC	GAAAAGAAC	AAAAGGCACC	80
JOR	80
ENG1	80
ENG/QA	80
MUN/ABD	80
FR1/UAE	80
FR2	56
	100	120	140	160					
EMC	AAAAGAAGAA	TCAACAGACCC	AAATGTCTGA	ACCTCCAAAG	GAGCAGCGTG	TGCAAGGTAG	CATCACTCGAG	CGCACTCGCA	160
JOR	160
ENG1	.	.	.	T	.	T	.	.	154
ENG/QA	160
MUN/ABD	160
FR1/UAE	160
FR2	136
	180	200	220	240					
EMC	CCCCGCCAAG	TGTTCAAGCCT	GGTCCAATGA	TTGATGTTAA	CACTGATTAG	TGTCACTCAA	AGTAACAAGA	TCGGGGCAAT	240
JOR	T	.	.	240
ENG1	240
ENG/QA	234
MUN/ABD	240
FR1/UAE	240
FR2	216
	260	280	300	320					
EMC	C	241							
JOR	.	241							
ENG1	.	241							
ENG/QA	.	235							
MUN/ABD	.	241							
FR1/UAE	.	241							
FR2	.	217							

Figure 1S: Alignments of the RdRp and N gene sequences the MERS-CoV of patients 1 and 2 with other available MERS-CoV sequences.

Sequences of patient 1 (FR1/UAE: Hu/France-UAE/1612/2013) and patient 2 (FR2: Hu/France/1631/2013) of the RdRp (nt 15073-15254) and N (nt 29598 or 29622-29838,) gene segments were aligned with available MERS-CoV sequences EMC: EMC/2012 (GenBank accession n°. JX869059), JOR: Jordan-N3/2012 (GenBank accession n°. KC776174), ENG1: England-1/2012 (GenBank accession n°. KC164505), ENG/QA: England/Qatar/2013 (GenBank accession n° KC667074), and MUN/ABD: Munich/Abu Dhabi/2013 (website Universitätsklinikum Bonn Institut für Virologie, <http://www.virology-bonn.de/index.php?id=46>; accessed on May 20, 2013). Dots represent identical nucleotides, hyphens represent sequence gaps.