# Evaluation of a SARS-CoV-2 rapid antigen test: potential to help reduce community spread?

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## 9 ABSTRACT

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Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) can spread from symptomatic 11 12 patients with COVID-19, but also from asymptomatic individuals. Therefore, robust surveillance 13 and timely interventions are essential for the control of virus spread within the community. In this 14 regard the frequency of testing and speed of reporting, but not the test sensitivity alone, play a 15 crucial role. In order to reduce the costs and meet the expanding demands in real-time RT-PCR 16 (rRT-PCR) testing for SARS-CoV-2, complementary assays, such as rapid antigen tests, have 17 been developed. Rigorous analysis under varying conditions is required to assess the clinical 18 performance of these tests and to ensure reproducible results. We evaluated the sensitivity and 19 specificity of a recently licensed rapid antigen test using 137 clinical samples in two institutions. 20 Test sensitivity was between 88.2-89.6% when applied to samples with viral loads typically seen 21 in infectious patients. Of 32 rRT-PCR positive samples, 19 demonstrated infectivity in cell culture, 22 and 84% of these samples were reactive with the antigen test. Seven full-genome sequenced 23 SARS-CoV-2 isolates and SARS-CoV-1 were detected with this antigen test, with no cross-24 reactivity against other common respiratory viruses. Numerous antigen tests are available for 25 SARS-CoV-2 testing and their performance to detect infectious individuals may vary. Head-to-

head comparison along with cell culture testing for infectivity may prove useful to identify better
performing antigen tests. The antigen test analyzed in this study is easy-to-use, inexpensive, and
scalable. It can be helpful in monitoring infection trends and thus has potential to reduce
transmission.

# 31 1. INTRODUCTION

Since the beginning of COVID-19 outbreak in December 2020, the global demand for the Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) testing has been steadily increasing. Already back in March 2020, hospitals and laboratories around the world announced their concerns about reagent, consumable material shortages, and limited personal protective equipment. Yet, timely detection and isolation of SARS-CoV-2 infected cases and identification of their contacts are pivotal to slowing down the pandemic.

38 The main public health strategy during a pandemic relies on robust and easy to perform 39 diagnostic tools that can be used to test large number of samples in a short time. To date the gold standard diagnostic method for SARS-CoV-2 detection [1] is based on real time reverse 40 transcription-PCR (rRT-PCR) technology which has been promptly implemented by the World 41 Health Organization (WHO) [2], Center for Disease Control and Prevention (CDC) [3] protocols, 42 and a number of commercial assays [4]. The SARS-CoV-2 rRT-PCR has high specificity and 43 44 sensitivity [5, 6]. However, the type and quality of the patient specimen [7, 8], stage of the disease, 45 and the degree of viral replication and/or clearance have an impact on the test outcome [9]. These 46 factors are critical not only for PCR-based but also for other diagnostic test systems aiming to detect the presence of the virus. Hence interpreting a test result for SARS-CoV-2 depends on the 47 accuracy of the test, but the prevalence and the estimated risk of disease before testing should 48 49 also be taken into consideration.

In many countries SARS-CoV-2 testing is extended to asymptomatic population, e.g. in schools, airports, nursing-homes, and workplaces. This leads to a growing gap between the large number of demand and the laboratory capacities to preform rRT-PCR tests, especially in developing countries. Despite high specificity and sensitivity, rRT-PCR has a disadvantage in point of care testing, because it usually requires professional expertise, expensive reagents and specialized equipment. Therefore, alternative assays, such as rapid antigen detection tests, which

56 can also detect the presence of the virus directly in respiratory samples, have been developed [4] 57 and tested by different groups [10-14]. However, it is vital to determine the sensitivity, specificity 58 of such tests relative to standard rRT-PCR in order to identify the ideal circumstances that their 59 application would be beneficial.

This study was performed to evaluate a novel antigen test produced by R-Biopharm for the detection of SARS-CoV-2 in different specimens and to identify its limitations and potential usage. Different types of materials and verification analysis were used by two institutions independently to assure the reproducibility of the testing and to analyze the potential caveats.

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#### 65 2. MATERIALS AND METHODS

#### 66 2.1 Specimen collection

At the Institute of Virology, Charité Berlin stored specimens taken after routine diagnostic were used with no extra procedures required for the study. Cell culture supernantants of respiratory viruses other than SARS-CoV-2 were available at the institute of virology, Charite through a EVD-LabNet EQA (https://www.evd-labnet.eu/; Fischer/Mögling, unpublished data).

At the Institute of Virology, Frankfurt, the clinical samples were collected from subjects as part of registered protocols. Combined oropharyngeal/nasal swabs were collected, stored in 2 ml PBS at 4°C and processed for further analysis within 24 hours.

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### 75 2.2 Cell culture and virus stocks

Caco-2 (human colon carcinoma) were cultured in Minimum Essential Medium (MEM) supplemented with 10% fetal calf serum (FCS). 100 IU/mL of penicillin and 100 g/mL of streptomycin. All culture reagents were purchased from Sigma (St. Louis. MO. USA). The Caco-2 cells were originally obtained from DSMZ (Braunschweig, Germany, no.: ACC 169) differentiated by serial passaging and selected for high permissiveness to virus infection. Caco-2 cells were

infected with different viral isolates (FFM1-FFM7) [15] at an MOI 0.1. Cell culture supernatant was
harvested 48 h after infection, precleared at 2000 x g for 10 min at room temperature. Aliquots of
virus particle containing supernatant were kept at -80°C.

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# 85 2.3 Detection of infectious virus in cell culture

Of the swab-dilution, 500 µL were mixed with 1.5 ml of MEM containing 1% FCS (Sigma-Aldrich;
St. Louis, Missouri, USA), 7.5 µg/ml Amphotericin B, and 0.1 mg/ml Primocin, (InvivoGen; San
Diego, California, USA). Swab-inoculums were transferred to Caco-2 cells seeded in 5.5 cm2
culture tubes. Cytopathogenic effect (CPE) was assessed daily for up to seven days or until cell
lysis occurred.

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# 92 2.4 Rapid Antigen Test

Rapid antigen test was provided by R-Biopharm. Test was performed according to the 93 94 manufacturer's recommendations and evaluated visually by four or six-eye principle. Briefly, 95 samples were vortexed for 20 sec. 50 µl from Solution A (blue) and B (yellow) were dispensed in clean 1.5 ml reaction tubes which leads to green coloring. Immediately 50 µl of the test samples 96 were added to the reaction mixture. Samples were then mixed briefly and incubated for 10 min at 97 98 room temperature. Test strips were placed in to mixture vertically to allow absorption. Test results 99 were evaluated after 10 min. Intensities of the test bands were compared to control band categorized as follows: +++ (test band intensity stronger than the control), ++ (test and control 100 bans intensity are similar), + (test band intensity is weaker than the control). Antigen testing for 101 viable SARS-CoV-2 and SARS-CoV-1 cell culture supernatants was performed in a BSL-3 102 103 laboratory.

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## 105 2.5 RNA extraction and rRT-PCR analysis

At the Institute of Virology, Charité Berlin, stored samples (swab resuspended in 1.5 mL of phosphate-buffered saline) were anonymized before testing. After thawing at RT all samples were analyzed by antigen test and rRT-PCR in parallel. RNA extraction for rRT-PCR was done by using the MagNA Pure 96 system, using 100 µl of sample, eluted in 100 µl. rRT-PCR was done as published previously [1].

At the Institute of Virology in Frankfurt the SARS-CoV-2 test (Cobas, Roche, Basel, Switzerland) was performed on the rRT-PCR automated Cobas 6800 system. Of the swab-dilution, 1000 µl aliquots were mixed with lysis buffer (1:1 ratio) and 500 µL aliquots were transferred to barcoded secondary tubes, loaded on the Cobas 6800 system, and tested with Cobas SARS-CoV-2 master mix containing an internal RNA control and primer-probe sets towards ORF1 and E-gene according to the manufacturer's instructions.

117 Within seven days of virus inoculation using clinical sample material, culture supernatant was

118 collected to perform rRT-PCR in order to confirm productive virus replication. RNA was isolated

119 from 100 µL cell culture supernatant using the QIAcube HT instrument and QIAamp 96 Virus

120 QIAcube HT Kit (Qiagen; Hilden, Germany) according to the manufacturer's instructions. SARS-

121 CoV-2 RNA was analyzed by rRT- PCR using the Luna Universal One-Step RT-qPCR Kit (New

122 England Biolabs; Ipswich, Massachusetts, USA) and primers targeting RNA-dependent RNA

polymerase (RdRp) [15]. RdRP\_SARSr-F2 (GTGARATGGTCATGTGTGGCGG)

124 RdRP\_SARSr-R1 (CARATGTTAAASACACTATTAGCATA).

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# 126 2.6 Statistical Analysis

The number of positive samples were compared two by two contingency table. The agreement between the antigen test and rRT-PCR techniques was evaluated using the Cohen's weighted kappa index (K value) [16]. K value interpretations were categorized as follows: <0.20 is poor, 0.21- 0.40 is fair, 0.41-0.60 is moderate agreement, 0.61- 0.80 is substantial agreement and 0.81-1.00 is almost perfect agreement [17].

# 132 2.7 Ethical Statement

The use of stored clinical samples for validation of diagnostic methods without person related data is covered by section 25 of the Berlin hospital law and does not require ethical or legal clearance. The use of anonymized clinical samples for validation of diagnostic methods does not require ethical clearance by the Goethe University, Frankfurt.

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# 138 3. RESULTS

Rapid antigen test sensitivity and specificity were evaluated by two independent institutions
using various number of clinical samples. rRT-PCR was used as a reference test system. We
deemed individuals to be uninfected with SARS-CoV-2 when a negative result was obtained by
rRT-PCR.



Figure 1. Antigen test analysis performed in Berlin (A) and Frankfurt (B). A. Log<sub>10</sub> RNA 144 copies/ml and corresponding antigen (Ag) detection test results (red circles positive n: 45, blue 145 146 circles negative n: 13) intensity for each rRT-PCR positive sample (n: 58). B. Cycle threshold (cT) value and corresponding antigen (Ag) detection test results (red circles positive n: 16, blue circles 147 negative n: 16) intensity for each rRT-PCR positive sample (n: 32). 32 rRT-PCR positive samples 148 were tested in cell culture for infectivity. All Ag-test positive (n:16, red circles) and three Ag-test 149 150 negative (red-filled blue circles) samples displayed CPEs after inoculating in Caco-2 cells (Table S2). Intensities of the test bands were compared to control band and designated as follows: +++ 151

(test band intensity stronger than the control), ++ (test and control bans intensity are similar), +
(test band intensity is weaker than the control).

In the Institute of Virology, Charité, Berlin, a total of 67 stored patient samples were 154 155 available for the study. Of these, 58 were rRT-PCR positive with cycle threshold (cT) range between 18.77-40 corresponding to 2.5x10<sup>9</sup> -1380 RNA genome copies/ml (Table S1), 156 representing 86.6% (58/67) of the clinical samples analyzed (Figure 1A). When the rRT-PCR 157 results were used as a reference, the antigen test diagnosed SARS-CoV-2 infection status with a 158 159 sensitivity of 77.6% (45/58) and a specificity of 100% (9/9) (Table 1). After re-evaluating the data based on the acceptable analytic sensitivity and limit of detection suggested by WHO [18], we 160 identified 48 samples with  $\geq 10^6$  RNA genome copies/ml. Rapid antigen test performed with 89.6% 161 sensitivity for this sample set (Table S1). Of these, 40 samples had ~2.23x10<sup>6</sup> or more RNA 162 genome copies/ml and reacted positive with the antigen test (Table 1). In contrast samples with 163 less than 7.63x10<sup>5</sup> RNA copies/ml were negative (Figure 1A, Table S1). Cohen's weighted kappa 164 165 value of 0.482 indicated moderate agreement between the rRT-PCR and the rapid antigen test 166 (Table 2). The overall concordance between the rRT-PCR and the antigen test was 80.6% (54/67).

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Antigen test							
		Negative	Positive	Marginal row	Sensitivity (%)	Specificity (%)	
rRT-PCR	Negative	9	0	9		100	
	Positive	13	45	58	77.6		
genome	≥2.23x10 <sup>6</sup>	0	40	40	100		
copies/ml	1.38x10 <sup>3</sup> -2.21x10 <sup>6</sup>	13	5	18	27.7%		

168

169 **Table 1**. Sensitivity and specificity of the antigen detection test in comparison to rRT-PCR

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	Antigen test		
rRT-PCR	Negative	Row marginal	
Negative	9	0	9 (13.4%)
Positive	13	45	58 (86.5%)
Column marginal	22 (32.8%)	67	

Weighted Kappa	0.482
Standard error	0.110
95% CI	0.266 to 0.698

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**Table 2**. Cohen's weighted kappa coefficient between rapid antigen test and rRT-PCR.

Certain rapid tests may be used at the point-of-care and thus offer benefits for the detection 174 175 and management of infectious diseases. In order to assess the potential of the rapid antigen test 176 in this context, 70 nasopharyngeal samples freshly collected from individuals living in a shared housing were analyzed head to head by rRT-PCR using Cobas 6800 system, rapid antigen test, 177 178 and cell culture using Caco-2 cells to determine the infectivity (Institute of Medical Virology, 179 Goethe University, Frankfurt). 45.7% (32/70) of the clinical samples were diagnosed positive for SARS-CoV-2 by rRT-PCR with cT values ranging between 18.01-35.98 (Figure 1B, Table S2). 180 181 The antigen test diagnosed the infection status with a sensitivity of 50% (16/32) and a specificity 182 of 100% (Table 3). Re-evaluating the data based on the limit of detection, sensitivity was 183 determined to be 88.2% for samples with cT values <28, and it was reduced in the group of 184 samples with cT values  $\geq$  28 (6.7%) (**Table 3**). Cohen's weighted kappa value of 0.521 indicated 185 moderate agreement between rRT-PCR and the rapid antigen test (Table 4). The overall concordance between the rRT-PCR and the antigen test was 77.1% (54/70) (Table 4). 186

187 rRT-PCR is a highly sensitive method to detect viral RNA molecules from clinical samples.
 188 However, viral RNA can persist in different body parts and can be detected in specimens for much
 189 longer than the presence of viable virus [19]. Thus demonstration of infectivity on permissive cell
 190 lines *in vitro* is a more reliable surrogate for infectivity and virus transmission. Therefore, we

191 attempted virus isolation by inoculating rRT-PCR positive samples in Caco-2 cells. Cytopathic changes were monitored daily by microscopy for a week and subsequently aliquots of culture 192 193 supernatant were tested to verify viral RNA copies (Table S2). For samples that are positive for 194 both antigen test and rRT-PCR (16/32, cT 18.01-28.45), we observed cytopathic effects (CPE) in cell culture 1-3 days after inoculation (Figure 1B, Table S2). Three samples that had a negative 195 result in the antigen test, but were positive by rRT-PCR (cT values 26.69, 30.12, and 32.13) 196 displayed CPE as well. Other 13 antigen-test negative samples with higher cT values (indicating 197 198 lower viral load) between 28.34-34.12 were not infectious in cell culture. Interestingly, one sample 199 with a relatively low cT value 25.53, did not show any CPE in cell culture and was also negative for the antigen test (Table S2). 200

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Antigen test						
Negative Positive Marginal Sensitivity Specificity row (%) (%)						
rRT-PCR	Negative	38	0	38		100
	Positive	16	16	32	50	
cT<28	Positive	2	15	17	88.2	
cT≥28	Positive	14	1	15	6.7	

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**Table 3**. Comparison of the clinical diagnostic performance of rapid antigen test with rRT-PCR.

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	Antigen test		
rRT-PCR	Negative	Row marginal	
Negative	38	0	38 (54.3%)
Positive	16 16		32 (45.7%)
Column marginal	54 (77.1%)	16 (22.9%)	70

Weighed Kappa	0.521
Standard error	0.092
95% CI	0.339 to 0.702

**Table 4**. Cohen's weighted kappa coefficient between rapid antigen test and rRT-PCR.

In order to investigate potential cross reactivity among common coronaviruses and other respiratory viruses, infectious and heat inactivated (4 h at 60°C) cell culture supernatants were tested (**Table 5**). SARS-CoV-1 and SARS-CoV-2 tested positive with the antigen test, as expected. The antigen test did not display any cross-reactivity with the other respiratory and endemic corona viruses listed in **Table 5**.

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Cell culture supernatant with virus	Ag-Test
SARS-CoV-1	+
SARS-CoV-2	+
HCoV-229E	-
HCoV-NL63	-
MERS	-
Enterovirus	-
Rhinovirus	-
Parainfluenzavirus 1	-
Parainfluenzavirus 2	-
Parainfluenzavirus 3	-
Parainfluenzavirus 4	-
hMPV A	-
hMPV B	-
RSV	-
Influenzavirus A H1N1	-
Influenzavirus A H3N2	-
Influenza B	-

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Table 5. Rapid Antigen Test results using different respiratory virus cell culture supernatantstocks.

We further evaluated the detection sensitivity among different SARS-CoV-2 isolates. Here we used cell culture supernatant collected from Caco-2 cells infected with seven different isolates [15] and SARS-CoV-1 (**Figure 3**). The virus stocks were thawed at room temperature and a total

219 of six 10-fold dilutions were prepared in PBS. The antigen test was performed and evaluated immediately (Figure 3A). In parallel, aliguots of the dilutions were mixed with lysis buffer used for 220 221 RNA extraction to inactivate the virus. rRT-PCR was performed for two different gene targets 222 ORF1 and E-gene that resulted in similar cT values (Figure 3B, Table S3). 10-fold serial dilutions 223 led to ~3 cT difference in rRT-PCR for each set as anticipated. According to our results the limit 224 of detection was between 100-560 RNA copies/ml which is in line with the manufacturer's findings. We previously identified RG203KR mutations in FFM3, FFM4 and FFM6 and S $\rightarrow$ L mutation in 225 FFM1 within the nucleocapsid protein coding region [15]. According to GISAID classification the 226 GR clade, carrying the combination of Spike D614G and nucleocapsid RG203KR mutations, is 227 228 currently the most common representative of the SARS-CoV-2 population worldwide [20]. Our 229 results suggest that the presence of the RG203KR mutation did not interfere with the antigen test 230 performance.



**Figure 3.** Rapid Antigen Test Results for SARS-CoV-1 and SARS-CoV-2 isolates. **A.** Representative lateral flow assay using serially diluted virus stock. Intensities of the test bands were compared to control band and designated as follows: +++ (test band intensity stronger than the control), ++ (test and control bans intensity are similar), + (test band intensity is weaker than the control). **B.** TCID50/ml values and corresponding antigen (Ag) detection test intensity for serially diluted SARS-CoV-2 isolates FFM1-7 and SARS-CoV-1 are shown. Representative result of two experiments.

## 240 4. DISCUSSION

241 In this study we validated the assay performance of a recently approved rapid antigen test 242 in two independent institutions using a total of 137 clinical samples. Although the test specificity was 100% for this particular sample set, overall sensitivity was low (50-77.6%), yet re-analyzing 243 244 samples with higher viral loads showed good correlation (88.2-89.6%). Previous studies reported 245 that lower cT values are associated with higher viral culture positivity [21, 22]. There is currently no direct evidence whether cell culture positivity or higher viral load correlates with contagiousness 246 247 of an individual, however, it is commonly recognized as the surrogate of infectivity [23]. Since an 248 important aspect of using point-of-care testing is to able to identify infected individuals who are infectious and can potentially transmit the virus, we performed correlation analysis within a group 249 of clinical samples tested. 19 out of 32 SARS-CoV-2 infected individuals were positive in cell 250 251 culture. The antigen test detected 16 out of 19 these (84%). In contrast 43.7% (14/32) of the 252 samples were not infectious in cell culture, yet positive by rRT-PCR, probably due to persisting 253 genomic and subgenomic viral RNA within the collected sample. We detected an excess amount 254 of viral RNA in cell culture supernatants due to high replication capacity of the virus in permissive 255 cells, despite a negative antigen test result. This might explain the cT discrepancy between the 256 cell culture supernatant and clinical samples. Limited clinical sample size is the major limitation of 257 this study. Future efforts should aim to monitor frequent sampling of larger groups and to compare different rapid antigen tests, different sampling sites along with infectivity correlation in cell culture. 258

Our results suggest that the rapid antigen test can detect SARS-CoV-2 infected individuals with high viral loads and has potential in determining highly contagious individuals. Despite low analytic sensitivity, rapid antigen tests are inexpensive and therefore can be used frequently for detecting infected individuals who are asymptomatic, pre-symptomatic and without known or suspected exposure to SARS-CoV-2 [24]. They can be beneficial in congregate settings, such as a long-term care facility or a correctional facility, workplace, or a school testing its students, faculty,

- and staff. Rapid antigen tests probably perform best during the early stages of SARS-CoV-2
- 266 infection when the viral load is higher.

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- heat inactivated respiratory viruses for specificity testing, Dr. Holger Rabenau for organizational
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# 280 SUPPLEMENTARY TABLES

- **Table S1.** Summary of rRT-PCR and antigen testing using clinical samples (Charite, Berlin). neg:
- 282 negative antigen test. N/D: not detectable.

Sample	rF	Antigen test	
ID	E- gene	Log10 RNA copies/mL	Intensity
#65	18,77	9,40	+++
#19	19,48	9,18	+++
#60	20,02	9,01	+++
#56	20,72	8,80	+++
#22	21,78	8,47	+++
#66	22,01	8,40	+++
#25	22,02	8,40	+++
#67	22,16	8,35	++
#21	22,74	8,17	+++
#20	22,94	8,11	+++
#57	23,11	8,06	++
#15	23,19	8,04	+++
#53	24,07	7,76	+++
#26	24,15	7,74	+++
#8	24,24	7,71	++
#12	25,15	7,43	++
#27	25,47	7,33	+
#58	25,7	7,26	+
#14	25,74	7,25	++
#1	25,92	7,19	++
#18	26,04	7,16	++
#64	26,12	7,13	+++
#4	26,14	7,13	+++
#10	26,22	7,10	++
#11	26,44	7,03	++
#23	26,71	6,95	++
#3	26,81	6,92	+++
#28	26,89	6,90	++
#29	26,89	6,90	++
#61	26,98	6,87	+++
#5	27,03	6,85	+
#13	27.05	6.85	++

1			
#59	27,09	6,83	+
#55	27,32	6,76	+
#49	27,47	6,72	+
#2	27,69	6,65	++
#17	27,85	6,60	++
#6	28,49	6,40	++
#7	28,52	6,39	++
#9	28,67	6,35	+
#46	28,68	6,34	neg
#50	28,92	6,27	neg
#51	29,21	6,18	+
#47	29,36	6,14	neg
#24	29,46	6,10	+
#32	29,59	6,06	+
#33	29,64	6,05	neg
#31	29,68	6,04	neg
#16	29,83	5,99	+
#54	29,87	5,98	+
#52	30,18	5,88	neg
#48	30,81	5,69	neg
#30	31,45	5,49	neg
#39	33,13	4,97	neg
#43	33,61	4,83	neg
#62	35,76	4,16	neg
#63	35,98	4,10	neg
#41	40	3,14	neg
#34	N/D	N/D	neg
#35	N/D	N/D	neg
#36	N/D	N/D	neg
#37	N/D	N/D	neg
#38	N/D	N/D	neg
#40	N/D	N/D	neg
#42	N/D	N/D	neg
#44	N/D	N/D	neg
#45	N/D	N/D	nea

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285

- 287 Table S2. Summary of rRT-PCR (Cobas), antigen testing, virus isolation in cell culture, rRT-PCR
- with cell culture supernatant of 70 clinical samples (University Hospital Frankfurt). neg: negative
- antigen test. N/D: not detectable.

Sample	Roche (COBAS) rRT-PCR		Antigen	CPE in cell	Luna One-step RdRP rRT-PCR	
ID	ORF1	E-gene	test	culture	supernatant	
1	18,01	18,01	+++	+	20,67	
2	20,22	20,42	+++	+	21,27	
3	21,45	21,31	++	+	20,69	
4	21,9	20,97	+	+	19,89	
5	21,96	22,27	+++	+	19,66	
6	22,2	22,39	+	+	20,72	
7	23,31	23,33	+	+	20,68	
8	23,33	23,21	++	+	20,41	
9	23,51	23,1	+	+	20,64	
10	23,76	23,7	+	+	20,22	
11	24,91	25,4	++	+	20,99	
12	25,5	20,00	neg	-	30,72	
14	20,4	20,03	nog	+	20,95	
15	26,30	26.2	+	+	20,32	
16	27.04	27.01	+	+	20,31	
17	27.46	27.11	+	+	20,23	
18	27,78	28,45	+	+	21,16	
19	28,39	28,92	neg	-	36,11	
20	28,8	28,34	neg	-	40,18	
21	29,32	30,12	neg	+	20,64	
22	29,84	30,54	neg	-	38,55	
23	30,15	31,03	neg	-	41,26	
24	30,45	31,35	neg	-	37,28	
25	31,49	32,77	neg	-	36,69	
26	31,89	32,99	neg	-	37,30	
27	32,19	33,06	neg	-	37,39	
28	32,34	34,03	neg	-	30,73	
29	35,09	32,13	neg	-	20,30	
31	35 39	33 67	neg	-	39.52	
32	35.78	34.12	nea	-	36.94	
33	36,3	N/D	neg	-	36,68	
34	N/D	N/D	neg		,	
35	N/D	N/D	neg			
36	N/D	N/D	neg			
37	N/D	N/D	neg			
38	N/D	N/D	neg			
39 40	N/D	N/D	neg			
41	N/D	N/D	nea			
42	N/D	N/D	neg			
43	N/D	N/D	neg			
44	N/D	N/D	neg			
45	N/D	N/D	neg			
46	N/D	N/D	neg			
47	N/D	N/D	neg			
48	N/D	N/D	neg			

49	N/D	N/D	neg
50	N/D	N/D	neg
51	N/D	N/D	neg
52	N/D	N/D	neg
53	N/D	N/D	neg
54	N/D	N/D	neg
55	N/D	N/D	neg
56	N/D	N/D	neg
57	N/D	N/D	neg
58	N/D	N/D	neg
59	N/D	N/D	neg
60	N/D	N/D	neg
61	N/D	N/D	neg
62	N/D	N/D	neg
63	N/D	N/D	neg
64	N/D	N/D	neg
65	N/D	N/D	neg
66	N/D	N/D	neg
67	N/D	N/D	neg
68	N/D	N/D	neg
69	N/D	N/D	neg
70	N/D	N/D	neq

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291

- **Table S3**. Cycle threshold (cT) value for ORF1 and E-gene rRT-PCR and rapid antigen test results
- for serially diluted SARS-CoV-2 isolates and SARS-CoV-1. neg: negative antigen test.

SAMPLE		rRT-PCR	(Cobas)	Ag-Test
		ORF1	E-gene	Intensity
	1,3E+05	13,12	13,86	+++
	1,3E+04	16,56	16,79	+++
SARS-CoV-2	1,3E+03	20,22	20,52	++
FFM1	1,3E+02	23,86	24,10	+
	1,3E+01	27,24	27,63	neg
	1,3E+00	30,50	31,02	neg
	1,0E+05	20,12	20,37	+++
	1,0E+04	23,23	23,62	++
SARS-CoV-2	1,0E+03	27,07	27,29	+
FFM2	1,0E+02	30,25	30,53	neg
	1,0E+01	33,24	33,83	neg
	1,0E+00	34,78	36,23	neg
	1,0E+05	16,36	16,63	+++
	1,0E+04	19,78	19,95	+++
SARS-CoV-2	1,0E+03	23,45	23,54	+
FFM3	1,0E+02	26,72	27,14	neg
	1,0E+01	30,15	30,68	neg
	1,0E+00	33,17	33,87	neg
	5,6E+05	12,7	13,4	+++
	5,6E+04	16,3	16,78	+++
SARS-CoV-2	5,6E+03	19,46	19,73	+++
FFM4	5,6E+02	23,3	23,5	+
	5,6E+01	26,61	26,93	neg
	5,6E+00	29,7	30	neg
	1,0E+05	13,96	14,08	+++
	1,0E+04	16,66	16,81	+++
SARS-CoV-2	1,0E+03	20,13	20,33	+++
FFM5	1,0E+02	23,26	23,64	++
	1,0E+01	27,01	27,08	+
	1,0E+00	29,74	30,11	neg
	1,0E+05	16,1	16,35	+++
	1,0E+04	19,57	19,83	+++
SARS-CoV-2	1,0E+03	23,43	23,47	++
FFM6	1,0E+02	26,36	26,63	+
	1,0E+01	30,04	30,64	neg
	1,0E+00	33,36	34,05	neg
	3,0E+05	14,87	15,12	+++
	3,0E+04	17,71	18,02	+++
SARS-CoV-2	3,0E+03	21,95	22,4	++
FFM7	3,0E+02	25,4	26,12	+
	3,0E+01	28,81	29,26	neg
	3,0E+00	32,11	32,72	neg
	1,0E+05	n/d	17	+++
	1,0E+04	n/d	20,2	++
SARS-CoV-1	1,0E+03	n/d	24,54	+
0, 110 000-1	1,0E+02	n/d	27,71	neg
	1,0E+01	n/d	31,1	neg
	1.0E+00	n/d	34.21	nea

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