



Analytical Sensitivity of the Abbott BinaxNOW COVID-19 Ag Card

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ABSTRACT Multiple rapid antigen (Ag) tests for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) have recently received emergency-use authorization (EUA) from the U.S. Food and Drug Administration (FDA). Although less sensitive than molecular detection methods, rapid antigen testing offers the potential for inexpensive, quick, decentralized testing. Robust analytical sensitivity data in comparison to reverse transcription-quantitative PCR (qRT-PCR) are currently lacking for many rapid antigen tests. Here, we evaluated the analytical sensitivity of the Abbott BinaxNOW COVID-19 Ag card using SARS-CoV-2-positive clinical specimens quantified by reverse transcription-droplet digital PCR (RT-ddPCR) and multiple FDA EUA qRT-PCR platforms using RNA standards. Initial and confirmatory limits of detection for the BinaxNOW COVID-19 Ag card were determined to be equivalent to 4.04×10^4 to 8.06×10^4 copies/swab. We further confirmed this limit of detection with 72 additional clinical samples positive for SARS-CoV-2 in either phosphate-buffered saline or viral transport medium. One hundred percent of samples with viral loads of $>40,000$ copies/swab were detected by rapid antigen testing. These data indicate that the BinaxNOW COVID-19 Ag card has an analytical sensitivity approximately equivalent to a generic qRT-PCR cycle threshold (C_T) value of 29 to 30.

KEYWORDS BinaxNOW, rapid antigen detection, SARS-CoV-2, limit of detection, coronavirus, COVID-19, Abbott, sensitivity

Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) continues to spread in the United States and across the world, causing tens of millions of cases and more than 1 million deaths. Testing for SARS-CoV-2 has predominantly been performed on molecular platforms with exquisite analytical sensitivities that can detect fewer than 100 viral RNA copies per ml of viral transport medium (VTM) (1). These analytical sensitivities result in the detection of low-level shedding in patients for weeks to months after initial infection, often with very low viral loads (2–5). A double-edged sword of the sensitivity offered by real-time reverse transcription-quantitative PCR (qRT-PCR) is the possibility that the molecular methodology detects RNA copies of virus that may not necessarily correlate with an active infection, especially if the result is a low viral load (6–8). However, assay sensitivity is critical for detecting additional cases where low viral loads in the nares are present during active lower respiratory tract infection (9).

Despite the current provision of more than 1 million SARS-CoV-2 tests per day in the United States, molecular testing may not be able to scale much further. Since the beginning of the pandemic, many have looked to antigen (Ag) tests to provide rapid, inexpensive, and decentralized testing that might potentially reduce transmission with the thought that antigen tests could successfully detect infectious cases (10, 11). However, the demand for antigen testing has outstripped the associated data on its performance, most notably with the \$750 million outlay by the U.S. government for

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the Abbott BinaxNOW COVID-19 Ag card based on data from just over 100 specimens in the emergency-use authorization (EUA) application (12–14). Only six samples with a cycle threshold (C_T) value of >33 were tested as part of the submission for U.S. Food and Drug Administration (FDA) authorization for the BinaxNOW COVID-19 Ag card. Other antigen tests have reported specificities of 100% based on testing tens of specimens, but real-world performance has not supported these estimates (15–17).

Accurately understanding the trade-off of sensitivity and speed is critical to achieving the right balance of diagnostics used in different settings. While determination of clinical sensitivity *in vivo* is the gold standard, such studies can be complicated by the ordering of swabs and by differences in patient anatomy, collector experience, and transport (18–20). Here, we evaluated the analytical sensitivity of the Abbott BinaxNOW COVID-19 Ag card in a central laboratory by subjecting the lateral flow assay to known amounts of SARS-CoV-2 quantified by a variety of qRT-PCR platforms.

MATERIALS AND METHODS

Specimen collection and quantification. This work was approved by the University of Washington institutional review board (IRB) (STUDY00010205). Deidentified samples used for the limit of detection (LoD) experiments were collected from patient specimens sent to the University of Washington Medical Center (UWMC) in Seattle, WA, for clinical diagnostic testing. Two nasopharyngeal swabs (NPSs) transported in phosphate-buffered saline (PBS) with high-viral-load C_T s of 16.0 and 21.9 by the Panther Fusion SARS-CoV-2 assay (Hologic, Marlborough, MA, USA) were pooled (21). This pool was quantified on the Roche cobas 6800 SARS-CoV-2 platform (Roche, Basel, Switzerland) using AccuPlex verification panel RNA standards (SeraCare Life Sciences, Milford, MA, USA) as well as by the reverse transcription-droplet digital PCR (RT-ddPCR) system (Bio-Rad, Hercules, CA, USA) using the CDC N1 primer (22, 23).

Four hundred microliters of input material was utilized for the Roche cobas SARS-CoV-2 assay, and 500 μ l was used for detection using the Hologic Panther Fusion SARS-CoV-2 assay. For RT-ddPCR, nucleic acids were extracted from 200 μ l of the sample using the QIAamp viral RNA minikit (Qiagen, Hilden, Germany) and eluted into 100 μ l distilled water (dH_2O); 10 μ l of this extracted RNA was used as the template in a 25- μ l amplification reaction mixture, with samples run in triplicate and evaluated by mean quantitation (23). The cobas 6800 calibration standards reported C_T s for the E gene of 27.72, 30.89, and 34.38 for 1×10^5 , 1×10^4 , and 1×10^3 calibration standards, respectively. The Panther Fusion assay standards reported C_T s for the open reading frame 1ab (ORF1ab) gene target of 27.3, 30.8, and 34.5 for 1×10^5 , 1×10^4 , and 1×10^3 calibrators, respectively. According to Hologic's FDA package insert, the Panther Fusion SARS-CoV-2 assay $1 \times$ LoD averaged from 10 contrived swab specimens was a C_T of 35.6, while the C_T cutoff is approximately 42 cycles (https://www.hologic.com/sites/default/files/2020-09/AW-21159-001_004_01.pdf). Similarly, Roche's FDA EUA reported the cobas 6800 LoD at mean C_T s of 32.7 for ORF1ab and 35.2 for the E gene, with a dynamic C_T cutoff of around 44 cycles, although it runs 50 cycles on the instrument (<https://www.fda.gov/media/136049/download>).

The Roche cobas assay, RT-ddPCR, and the Abbott BinaxNOW rapid antigen test were all performed in parallel. Additionally, BinaxNOW provided positive swabs that were tested for each shipment lot number to account for reagent and Ag assay quality control. For analytical negative controls, at both initial and confirmatory LoD dilutions, a clinical NPS previously determined to be negative by the Panther Fusion assay was subjected to the rapid Ag test.

Determination of analytical sensitivity. For the initial LoD, rapid Ag tests were run in triplicate with serial 10-fold PBS dilutions of the virus. First, a range of volumes (dH_2O) from 25 to 100 μ l was applied to each kit-provided sterile foam-tipped applicator to visually evaluate saturation; after soaking the swab in dH_2O , any liquid remaining in the microcentrifuge tube was measured with a pipette. Without oversaturation or underexposing the swab, 50 μ l was determined to be appropriate for saturation. Next, 50 μ l of the quantified sample was aliquoted into microcentrifuge tubes, where each swab was inserted, gently rotated to absorb liquid, and subjected to the rapid antigen test according to the manufacturer's protocol.

To account for liquid absorbed by the swab that might dilute the extraction buffer, a separate set of triplicate swabs was also allowed to air dry for 25 min after being saturated. To account for the potential impact of PBS on the antigen assay reaction, the original pool was also diluted in triplicate with dH_2O for comparison. No difference was detected in either comparison for dH_2O -diluted swab saturation and air-dried swabs. The initial LoD in triplicate was determined at the same dilution level regardless of whether the virus applied to the swab was diluted in PBS, diluted in water, or air dried (data not shown). All COVID-19 Ag cards were read after 15 min, and samples were never frozen and stored at 4°C.

RESULTS

The original positive sample pool yielded C_T s of 17.26 for target 1 (ORF1ab) and 17.27 for target 2 (E gene) with the Roche cobas SARS-CoV-2 assay. Based on RNA standard quantification using the E gene primer set, this result corresponded to

1.13×10^8 copies/ml, while RT-ddPCR quantified the sample at 5.65×10^7 copies/ml based on the CDC N1 primer set. Initial LoDs of BinaxNOW were recognized as the last dilution set to detect 3/3 (100%) positive samples. At first, this put the detection range between 10-fold and 100-fold dilutions of the neat, positive pool (or 2.83×10^5 to 2.83×10^4 copies/swab when quantified by RT-ddPCR) (Table 1).

Further dilution sets in triplicate narrowed this initial LoD range to 1:70 to 1:80 dilutions. Confirmatory LoDs were established with 20 replicates at each 1:70 and 1:80 dilution and characterized by $\geq 95\%$ (19/20) positive replicates detected. Initial and confirmatory LoDs for Abbott's BinaxNOW rapid antigen test were both determined at a 1:70 dilution of the original pooled positive samples, corresponding to 8.06×10^4 E gene copies/swab by RNA standards run on Roche cobas 6800 or 4.04×10^4 N gene copies/swab by RT-ddPCR (Table 1).

Based on the RNA standard curve on the cobas 6800 platform and a typical dilution of a swab into 3 ml VTM or PBS, these limits of detection correspond to E gene C_T s of approximately 30.5 and 29.5, respectively. We then took the 50- μ l volume at the LoD (1:70 dilution of the neat, positive pool), applied it to the kit-provided foam-tipped swab, and diluted it in 3 ml of PBS, similar to how a clinical NPS would be treated for molecular detection. The cobas assay quantified this sample with C_T s of 28.68 (ORF1ab) and 29.08 (E gene).

After establishing the LoD range at 4.04×10^4 to 8.06×10^4 copies/swab, we next expanded the sample set to include 72 additional clinical positive specimens. These fresh NPSs were previously positive by Hologic Panther Fusion with ORF1ab C_T s ranging from 14.6 to 39.9, corresponding to viral loads of 3.29×10^8 to 2.91×10^1 copies/ml, and stored in either PBS or VTM. For all specimens with $>40,000$ copies/swab ($n=24$), the Abbott BinaxNOW-19 Ag card correctly detected 100% of deidentified samples (Table 2). Notably, multiple samples beyond the LoD were still detected, with some having as few as an estimated 2,400 copies/swab.

DISCUSSION

Here, we describe a detailed examination of the analytical sensitivity of the BinaxNOW COVID-19 Ag card in a central laboratory. We estimate a limit of detection of 4.04×10^4 to 8.06×10^4 viral copies/swab, corresponding to a C_T of approximately 30. These data generally agree with Abbott's package insert, which separated samples based on a C_T cutoff of 33. These data suggest a difference of approximately 6 to 7 C_T s between the LoD of this antigen test and that of RT-PCR tests, indicating an ~ 100 -fold difference in sensitivity. Importantly, the difference in the C_T is notable as several studies have been able to culture virus at viral loads above a C_T of 30 but rarely above a C_T of 34 (8, 24, 25). According to work by Singanayagam et al., viable viruses with C_T s of >35 were found in 8.3% (5/60) of patient samples (26). Importantly, putting the limit of detection in terms of qRT-PCR C_T s can vary by platform based on extraction and elution volumes, viral transcripts targeted by the assay, and the amount of transport medium that the swab is diluted in before it is amplified for molecular detection. To avoid being dependent on one assay for quantitation, we quantified our original pooled positive samples in PBS on multiple platforms using multiple quantitation methods in our clinical laboratory in terms of viral copies per swab. These values of copies per swab correlate the amount of virus that each rapid antigen test sees with molecular detection methods.

This study was chiefly limited by the testing of specimens sent in transport media to a central laboratory. Rapid antigen tests are not generally meant to be tested on specimens in transport media, overcoming their higher limits of detection by sampling a whole swab. We took measures to ensure that our contrived specimens would directly answer the question of limits of detection by first using specimens in PBS for the LoD determination since the compositions of viral transport media and other complex matrices have been shown to inhibit rapid antigen assays (14, 27). We focused on rigorously interrogating the analytical sensitivity of the assay and did not evaluate

TABLE 1 Initial and confirmatory LoDs for the Abbott BinaxNOW COVID-19 Ag card^b

Sample	Dilution	No. of initial replicates detected/ no. tested	No. of confirmatory replicates detected/ no. tested	Initial % positive	Confirmatory % positive	No. of copies/ml (ddPCR)	No. of copies/swab (ddPCR)	No. of copies/ml (cobas)	No. of copies/swab (cobas)
S0 ^a	Neat	3/3		100		5.65 × 10 ⁷	2.83 × 10 ⁶	1.13 × 10 ⁸	5.64 × 10 ⁶
S1	1:10	3/3		100		5.65 × 10 ⁶	2.83 × 10 ⁵	1.13 × 10 ⁷	5.64 × 10 ⁵
S1-2	1:20	3/3		100		2.83 × 10 ⁶	1.42 × 10 ⁵	5.64 × 10 ⁶	2.82 × 10 ⁵
S1-4	1:40	3/3		100		1.41 × 10 ⁶	7.05 × 10 ⁴	2.82 × 10 ⁶	1.41 × 10 ⁵
S1-5	1:50	3/3		100		1.13 × 10 ⁶	5.65 × 10 ⁴	2.26 × 10 ⁶	1.13 × 10 ⁵
S1-6	1:60	3/3		100		9.42 × 10 ⁵	4.71 × 10 ⁴	1.88 × 10 ⁶	9.40 × 10 ⁴
S1-7	1:70	3/3	20/20	100	100	8.07 × 10⁵	4.04 × 10⁴	1.61 × 10⁶	8.06 × 10⁴
S1-8	1:80	3/3	17/20	67	85	7.06 × 10⁵	3.53 × 10⁴	1.41 × 10⁶	7.05 × 10⁴
S2	1:1,000	0/3		0		5.65 × 10 ⁵	2.83 × 10 ⁴	1.13 × 10 ⁶	5.64 × 10 ⁴
S3	1:1,000	0/3		0		5.65 × 10 ⁴	2.83 × 10 ³	1.13 × 10 ⁵	5.64 × 10 ³
Positive control	Neat	1/1	1/1	100	100	NA	NA	NA	NA
Negative control	Neat	0/1	0/1	0	0	NA	NA	NA	NA

^aS0, the undiluted positive pool, was determined to have a C_t of 17.3 for both targets on the Roche cobas 6800 platform.

^bThe positive control was a kit-provided swab by Abbott; the negative control was a clinical nasopharyngeal swab previously determined to be negative by Hologic Panther Fusion. Abbreviations: ddPCR, droplet digital PCR; Pos, positive; Neg, negative; NA, not applicable. The two dilutions that confirm the LoD are in boldface.

TABLE 2 COVID-19 Ag card rapid test across a range of SARS-CoV-2-positive specimens^a

UW sample ID	Matrix medium	Panther Fusion C _T	Original no. of copies/ml	No. of copies/swab	No. of copies/ml after dilution in 3 ml	Theoretical Panther Fusion C _T	BinaxNOW antigen detection
8	VTM	14.6	3.29 × 10 ⁸	1.65 × 10 ⁷	5.49 × 10 ⁶	21.0	DET
3	PBS	17.1	6.65 × 10 ⁷	3.33 × 10 ⁶	1.11 × 10 ⁶	23.5	DET
13	VTM	17.1	6.65 × 10 ⁷	3.33 × 10 ⁶	1.11 × 10 ⁶	23.5	DET
9	VTM	17.4	5.49 × 10 ⁷	2.75 × 10 ⁶	9.15 × 10 ⁵	23.8	DET
72	VTM	17.6	4.83 × 10 ⁷	2.42 × 10 ⁶	8.06 × 10 ⁵	24.0	DET
67	VTM	17.8	4.25 × 10 ⁷	2.13 × 10 ⁶	7.09 × 10 ⁵	24.2	DET
19	VTM	18	3.74 × 10 ⁷	1.87 × 10 ⁶	6.24 × 10 ⁵	24.4	DET
54	VTM	18	3.74 × 10 ⁷	1.87 × 10 ⁶	6.24 × 10 ⁵	24.4	DET
71	VTM	18.1	3.51 × 10 ⁷	1.76 × 10 ⁶	5.85 × 10 ⁵	24.5	DET
60	VTM	18.2	3.29 × 10 ⁷	1.65 × 10 ⁶	5.49 × 10 ⁵	24.6	DET
16	VTM	18.5	2.72 × 10 ⁷	1.36 × 10 ⁶	4.53 × 10 ⁵	24.9	DET
44	VTM	18.7	2.39 × 10 ⁷	1.20 × 10 ⁶	3.99 × 10 ⁵	25.1	DET
42	VTM	19.3	1.63 × 10 ⁷	8.15 × 10 ⁵	2.72 × 10 ⁵	25.7	DET
47	VTM	19.5	1.43 × 10 ⁷	7.17 × 10 ⁵	2.39 × 10 ⁵	25.9	DET
64	VTM	19.7	1.26 × 10 ⁷	6.31 × 10 ⁵	2.10 × 10 ⁵	26.1	DET
6	PBS	19.9	1.11 × 10 ⁷	5.55 × 10 ⁵	1.85 × 10 ⁵	26.3	DET
10	VTM	19.9	1.11 × 10 ⁷	5.55 × 10 ⁵	1.85 × 10 ⁵	26.3	DET
69	VTM	21	5.50 × 10 ⁶	2.75 × 10 ⁵	9.16 × 10 ⁴	27.4	DET
17	VTM	21.2	4.84 × 10 ⁶	2.42 × 10 ⁵	8.06 × 10 ⁴	27.6	DET
63	VTM	21.2	4.84 × 10 ⁶	2.42 × 10 ⁵	8.06 × 10 ⁴	27.6	DET
48	VTM	21.5	3.99 × 10 ⁶	2.00 × 10 ⁵	6.65 × 10 ⁴	27.9	DET
61	VTM	22	2.90 × 10 ⁶	1.45 × 10 ⁵	4.83 × 10 ⁴	28.4	DET
12	VTM	22.1	2.72 × 10 ⁶	1.36 × 10 ⁵	4.53 × 10 ⁴	28.5	DET
52	VTM	22.1	2.72 × 10 ⁶	1.36 × 10 ⁵	4.53 × 10 ⁴	28.5	DET
70	VTM	22.5	2.11 × 10 ⁶	1.05 × 10 ⁵	3.51 × 10 ⁴	28.9	DET
22	VTM	23.1	1.44 × 10 ⁶	7.18 × 10 ⁴	2.39 × 10 ⁴	29.5	DET
51	VTM	23.4	1.18 × 10 ⁶	5.92 × 10 ⁴	1.97 × 10 ⁴	29.8	DET
49	VTM	23.8	9.17 × 10 ⁵	4.59 × 10 ⁴	1.53 × 10 ⁴	30.2	NOT DET
58	VTM	23.9	8.60 × 10 ⁵	4.30 × 10 ⁴	1.43 × 10 ⁴	30.3	NOT DET
65	VTM	24.2	7.10 × 10 ⁵	3.55 × 10 ⁴	1.18 × 10 ⁴	30.6	DET
57	VTM	24.4	6.25 × 10 ⁵	3.12 × 10 ⁴	1.04 × 10 ⁴	30.8	DET
32	VTM	24.7	5.16 × 10 ⁵	2.58 × 10 ⁴	8.60 × 10 ³	31.1	NOT DET
23	VTM	24.8	4.84 × 10 ⁵	2.42 × 10 ⁴	8.06 × 10 ³	31.2	NOT DET
1	PBS	25	4.26 × 10 ⁵	2.13 × 10 ⁴	7.10 × 10 ³	31.4	DET
68	VTM	25.1	3.99 × 10 ⁵	2.00 × 10 ⁴	6.66 × 10 ³	31.5	NOT DET
43	VTM	25.4	3.30 × 10 ⁵	1.65 × 10 ⁴	5.50 × 10 ³	31.8	NOT DET
59	VTM	25.4	3.30 × 10 ⁵	1.65 × 10 ⁴	5.50 × 10 ³	31.8	DET
29	VTM	26.9	1.26 × 10 ⁵	6.32 × 10 ³	2.11 × 10 ³	33.3	NOT DET
55	VTM	27.1	1.11 × 10 ⁵	5.56 × 10 ³	1.85 × 10 ³	33.5	DET
27	VTM	27.2	1.04 × 10 ⁵	5.21 × 10 ³	1.74 × 10 ³	33.6	DET
33	VTM	27.4	9.18 × 10 ⁴	4.59 × 10 ³	1.53 × 10 ³	33.8	NOT DET
35	VTM	27.7	7.58 × 10 ⁴	3.79 × 10 ³	1.26 × 10 ³	34.1	DET
38	VTM	27.7	7.58 × 10 ⁴	3.79 × 10 ³	1.26 × 10 ³	34.1	DET
21	VTM	28.1	5.87 × 10 ⁴	2.93 × 10 ³	9.78 × 10 ²	34.5	NOT DET
37	VTM	28.1	5.87 × 10 ⁴	2.93 × 10 ³	9.78 × 10 ²	34.5	DET
62	VTM	28.1	5.87 × 10 ⁴	2.93 × 10 ³	9.78 × 10 ²	34.5	NOT DET
53	VTM	28.3	5.16 × 10 ⁴	2.58 × 10 ³	8.60 × 10 ²	34.7	NOT DET
2	PBS	28.4	4.84 × 10 ⁴	2.42 × 10 ³	8.07 × 10 ²	34.8	DET
46	VTM	28.4	4.84 × 10 ⁴	2.42 × 10 ³	8.07 × 10 ²	34.8	NOT DET
45	VTM	28.6	4.26 × 10 ⁴	2.13 × 10 ³	7.10 × 10 ²	35.0	NOT DET
11	VTM	29	3.30 × 10 ⁴	1.65 × 10 ³	5.50 × 10 ²	35.4	NOT DET
26	VTM	29	3.30 × 10 ⁴	1.65 × 10 ³	5.50 × 10 ²	35.4	NOT DET
41	VTM	29	3.30 × 10 ⁴	1.65 × 10 ³	5.50 × 10 ²	35.4	NOT DET
34	VTM	29.1	3.09 × 10 ⁴	1.55 × 10 ³	5.16 × 10 ²	35.5	NOT DET
66	VTM	29.4	2.55 × 10 ⁴	1.28 × 10 ³	4.26 × 10 ²	35.8	NOT DET
56	VTM	30.1	1.63 × 10 ⁴	8.16 × 10 ²	2.72 × 10 ²	36.5	NOT DET
18	VTM	30.6	1.19 × 10 ⁴	5.93 × 10 ²	1.98 × 10 ²	37.0	NOT DET
50	VTM	30.8	1.04 × 10 ⁴	5.22 × 10 ²	1.74 × 10 ²	37.2	NOT DET
39	VTM	31	9.18 × 10 ³	4.59 × 10 ²	1.53 × 10 ²	37.4	NOT DET
36	VTM	31.3	7.58 × 10 ³	3.79 × 10 ²	1.26 × 10 ²	37.7	NOT DET
25	VTM	31.6	6.26 × 10 ³	3.13 × 10 ²	1.04 × 10 ²	38.0	NOT DET

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TABLE 2 (Continued)

UW sample ID	Matrix medium	Panther Fusion C_T	Original no. of copies/ml	No. of copies/swab	No. of copies/ml after dilution in 3 ml	Theoretical Panther Fusion C_T	BinaxNOW antigen detection
14	VTM	31.7	5.87×10^3	2.93×10^2	9.78×10^1	38.1	NOT DET
24	VTM	32.1	4.54×10^3	2.27×10^2	7.57×10^1	38.5	NOT DET
30	VTM	34.1	1.26×10^3	6.32×10^1	2.11×10^1	40.5	NOT DET
20	VTM	35	7.11×10^2	3.56×10^1	1.19×10^1	>40.5	NOT DET
31	VTM	35.1	6.67×10^2	3.34×10^1	1.11×10^1	>40.5	NOT DET
4	PBS	36.2	3.30×10^2	1.65×10^1	5.50	>40.5	NOT DET
28	VTM	36.4	2.91×10^2	1.45×10^1	4.84	>40.5	NOT DET
40	VTM	36.8	2.25×10^2	1.13×10^1	3.75	>40.5	NOT DET
15	VTM	38.1	9.80×10^1	4.9	1.63	>40.5	NOT DET
5	PBS	38.8	6.26×10^1	3.13	1.04	>40.5	NOT DET
7	PBS	39.9	3.10×10^1	1.55	0.52	>40.5	NOT DET

^aAbbreviations: Ag, antigen; UW, University of Washington; C_T , cycle threshold; VTM, viral transport medium; PBS, phosphate-buffered saline; DET, detected; NOT DET, not detected. The conversion factor for copies per swab is 1/20 of copies per milliliter. Theoretical Panther Fusion C_T s were calculated with RNA standards factoring in a 1:20 dilution for the copies per swab and then an additional dilution in 3 ml of transport medium. Hologic Panther Fusion C_T s of >40.5 are expected to be missed.

assay specificity as we presumed that specimens in transport media would not adequately reflect relevant interferences found in direct nasal specimens. It has been well established that molecular detection is not grossly affected by PBS versus VTM (28). Moreover, two studies have used PBS or VTM samples for antigen testing (13, 29). It is possible that additional studies could better inform the equivalency of VTM versus PBS. We also note that the assays performed here do not adjudicate the “infectiousness” of the individuals associated with these specimens or the public health impact of different testing algorithms, which can be measured only by detailed clinical investigation outside the laboratory.

Our work also did not measure the analytical specificity of the antigen test as we believed that we could not adequately model specificity under the contrived conditions in the laboratory. Understanding assay specificity is critical to estimating the number of false-positive results that may occur in a given testing environment. Of note, the FDA recently administered a press release on 3 November 2020, warning clinical laboratory staff and health care providers to be aware of the potential for false-positive test results with rapid antigen tests (30). Finally, in order to compare sensitivities in terms of qRT-PCR, we reported the LoD in viral RNA copies, even though viral RNA is not targeted by rapid antigen tests. Although C_T s can vary between platforms, laboratories, analytical parameters, and even technicians and runs, LoD C_T values are also reported here for semiquantitative reference.

Having access to affordable, rapid testing is critical to reducing transmission during a pandemic. Although rapid antigen tests can be convenient and inexpensive, here, we confirm that thousands to tens of thousands of viral copies are necessary for detection, which is significantly greater than that required for qRT-PCR (12, 13, 31). These limitations in sensitivity are balanced by the ease of use and short turnaround time of the antigen test (32, 33). Our work more closely approximates the inherent abilities and limitations of antigen cards to detect virus, beyond the myriad of preanalytical variables that may differ between tests. These estimates are critical for evaluating when to use antigen versus qRT-PCR testing and to more accurately model how many cases are detected and missed by antigen tests (34). A limit of detection at ~40,000 copies/swab was also recently seen in outside work performed in parallel (35). More epidemiological and clinical research work, beyond viral culture, is required to determine whether the viral loads detected by antigen cards correspond to “infectious” virus and whether rapid diagnostics will ultimately help to detect and reduce viral transmission.

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