Relationship of Viral Load and Infectivity to the Limit of Detection of SARS-CoV-2 Antigen Tests

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OUESTION: When/how should we use SARS-CoV-2 antigen versus PCR tests?

ANTIGEN TESTS

RAPID INEXPENSIVE POINT-OF-CARE INSENSITIVE?

> **PCR** SLOW EXPENSIVE SENSITIVE

INFECTION VERSUS INFECTIVITY INFECTIVITY SURROGATE = VIRAL CULTURE

log10 day 3 culture upernatant viral load Sensitivity vs. LoD 1.0 -0.9 0.8 0.7 Abbott M2000 EUA CDC and GenMark EUA - LabCorp EUA S 0.6 0.5 € 0.4 Sofia2 Ag Test EUA 0.3 0.2 0.1

From Arnaout et al. CID PMC7302192

 $10^{1}10^{2}10^{3}10^{4}10^{5}10^{6}10^{7}10^{8}10^{9}$

Genome copies/mL

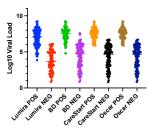


Figure 1. Antigen Testing Results Compared with Log10 Viral Load. Viral load in genome copies/mL POS = positive antigen test result. NEG = negative antigen testing rest. Lumira = LumiraDx Aq test: BD = BD Veritor Aq test.

Sensitivity/Specificity versus Viral Culture LumiraDx 90% (83-94% C.I.) / 70% (59-79%) Other Ag 74% (65-82% C.I.) / 92% (84-96%)

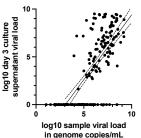
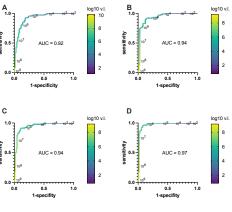
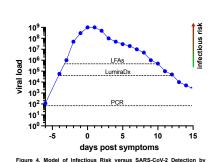


Figure 2. Quantitative Relationship Between Culturable Virus and Sample Viral Load. Day 3 viral culture supernatant for each sample was analyzed by RT-qPCR. The viral load in log10 genome copies/mL of culture supernatant is plotted against the log10 viral load in genome copies/mL of the original patient sample. Linear regression (solid line) with 95% confidence intervals (dashed lines) shown, R2 = 0.55



risk to others. Dotted lines indicate reliable detection threshold predicted for each method. Presumptively, infectious risk is proportional to the amount of culturable virus which is roughly proportional to the viral load in samples. Figure 3. Receiver operator curves (ROC) comparing Antigen tests are excellent in detecting patients with the highest viral loads SARS-CoV-2 sample viral load levels as a predictor of which may be four to five log10-fold greater than viral loads detected at the viral culture and antigen detection. For each plot, lowest levels where virus can be consistently cultured. PCR and to a lesser sensitivity versus 1-specificity was plotted for each viral extent, the LumiraDx test, can detect individuals before and after the expected load value (genome/copies/mL) determined by RT-gPCR for each sample in our study when used as a lower limit infectious period and therefore may be more appropriate for screening threshold for scoring positive and negative detection for all programs where testing is performed at longer intervals. The viral load curve other viral load results with qualitative viral culture or shown is for representational purposes and may not reflect viral load kinetics in antigen test determinations, respectively, as the any specific individual. comparators. (A) Log10 viral load (v.l.) in genome copies/ mL versus detection by viral culture. (B) Log10 viral load versus LumiraDx antigen detection. (C) Log10 viral load versus BD Veritor antigen detection. (D) Log10 viral load versus Oscar Biosciences antigen detection. (E) Log10 viral load versus CareStart antigen detection. Viral load values along the ROC curves are labeled in log10 intervals



RT-qPCR and Antigen Tests. Both Lumira and lateral flow-based antigen tests

(e.g., BD Veritor, CareStart, and Oscar Biosciences) are able to detect individu-

als with viable, culturable virus and who therefore pose an immediate infectious

0.5 1-specifity Conclusions:

0.0

AUC = 0.97

- Use Ag tests to identify *infectious* individuals at time of testing. Will allow isolation of significantly infectious individuals from communal events, same-day healthcare procedures, communal travel arrangements, and other functions with significant person-to-person contact in settings where universal masking is neither feasible nor desired.
- PCR tests for no-margin-for-error situations (hospital admission), vulnerable populations; sample pooling strategies; and screening of cohorted populations (e.g., school) at decreased intervals.

and demarcated in color as indicated in accompany

heatmap legend bar. AUC (area under the curve) for each ROC curve is denoted on respective plots.

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