

# Evaluation of Severe Acute Respiratory Syndrome Coronavirus 2 Nucleocapsid Antigen in the Blood as a Diagnostic Test for Infection and Infectious Viral Shedding

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**Background.** SARS-CoV-2 nucleocapsid antigen can be detected in plasma, but little is known about its performance as a diagnostic test for acute SARS-CoV-2 infection or infectious viral shedding among nonhospitalized individuals.

*Methods.* We used data generated from anterior nasal and blood samples collected in a longitudinal household cohort of SARS-CoV-2 cases and contacts. Participants were classified as true positives if polymerase chain reaction (PCR) positive for SARS-CoV-2 and as true negatives if PCR negative and seronegative. Infectious viral shedding was determined by the cytopathic effect from viral culture. Stratified by 7 days after symptom onset, we constructed receiver operating characteristic (ROC) curves to describe optimized accuracy (Youden index), optimized sensitivity, and specificity.

**Results.** Of 80 participants, 58 (73%) were true positives while 22 (27%) were true negatives. Using the manufacturer's cutoff of 1.25 pg/mL for evaluating infection, sensitivity was higher from 0 to 7 days (77.6% [95% confidence interval {CI}, 64%–88.2%]) than from 8 to 14 days (43.2% [95% CI, 31.1%–54.5%]) after symptom onset; specificity was unchanged at 100% (95% CI, 88.1%–100%). This test had higher sensitivity (100% [95% CI, 88.4%–100%]) and lower specificity (65% [95% CI, 40.8%–84.6%]) for infectious viral shedding as compared with infection, particularly within the first week of symptom onset. Although the presence of N-antigen correlated with infectious viral shedding (r = 0.63; P < .01), sensitivity still declined over time. Additional cutoffs from ROC curves were identified to optimize sensitivity and specificity.

**Conclusions.** We found that this SARS-CoV-2 N-antigen test was highly sensitive for detecting early but not late infectious viral shedding, making it a viable screening test for community-dwelling individuals to inform isolation practices.

Keywords. blood; infectiousness; infectivity; nucleocapsid antigen; performance.

As of April 2022, >847 million severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) tests have been performed in the United States [1], resulting in >80 million infections diagnosed. Screening and diagnostic tests have been developed to detect SARS-CoV-2 RNA, antibodies, or antigens [2–4]. Nucleocapsid protein (N-antigen) was identified as one of the predominantly expressed proteins that could have comparable performance with nucleic acid amplification tests, particularly

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early in the course of disease [5]. Although antigen testing is generally performed with nasal and saliva biospecimens, emerging evidence suggests that detecting viral proteins in plasma might represent a novel approach to improve the screening or diagnosis of SARS-CoV-2 infection [6–9], depending on its test performance characteristics in various populations.

Although almost all reports of N-antigen in the plasma involve characterization of hospitalized individuals [10, 11], there are few reports among mildly ill or asymptomatic individuals, groups that constitute the majority of individuals infected with SARS-CoV-2. One study comparing the single-molecule array (Simoa) SARS-CoV-2 N-antigen assay (Quanterix, Billerica, Massachusetts) [12] in plasma samples using a cutoff of 1.25 pg/ mL with nasal RNA positivity evaluated test performance characteristics among long-term care and hospitalized individuals; the authors found 100% specificity and 97.5% sensitivity [13]. This study

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did not include individuals living in the community, an important population with potentially different performance characteristics.

There has been a growing evidence base that nasal and saliva antigen testing can provide some indication about infectiousness, but these estimates are imprecise. The challenge with routine molecular testing for SARS-CoV-2 is that it cannot differentiate between the active virus replication (infectious virus) from persistent viral shedding (noninfectious virus). Nasal and saliva antigen testing can provide some indication about infectiousness, but these estimates are imprecise and no single laboratory test has been able to serve as a reliable predictor of infectious viral shedding [14–17]. As there is an urgent need to identify a test to predict infectiousness, we must continue to explore how novel diagnostic approaches may improve our ability to predict infectious viral shedding from blood-based N-antigen tests.

Utilizing a nonhospitalized cohort of individuals infected with SARS-CoV-2 and their household contacts [18], we aimed to evaluate the sensitivity and specificity of the SARS-CoV-2 N-antigen test on plasma (Simoa technology) as compared to infection, stratifying by time since symptom onset. We further investigated the presence and magnitude of SARS-CoV-2 N-antigen in plasma over time and its relationship with the levels and duration of infectious virus measured in nasal samples.

#### METHODS

#### **Patient Consent Statement**

The study protocol was reviewed by the University of California, San Francisco (UCSF) and Centers for Disease Control and Prevention institutional review boards (IRBs) and was determined to meet criteria for public health surveillance activities and did not require IRB oversight, according to the federal regulations summarized in 45 Code of Federal Regulations 46.102(1)(2). Written informed consent was obtained from all participants.

#### Study Design, Participants, and Procedures

We used data generated from a subgroup of participants enrolled in our natural history study of acute SARS-CoV-2 infectivity [18]. The parent study was an observational longitudinal cohort based in the San Francisco Bay Area, which has been previously described in detail [19]. In brief, index cases with  $\geq$ 1 household member were identified from SARS-CoV-2 RNA-positive individuals at UCSF-affiliated health facilities; households were enrolled within 5 days of symptom onset. Interview-administered questionnaires and blood samples were obtained at enrollment, and at days 9, 14, 21, and 28 after the index case's symptom onset. Anterior nasal specimens were self-collected daily through day 14 and on days 17, 19, 21, and 28. For this analysis, we used the plasma and nasal biospecimens collected from consecutive participants enrolled into the parent cohort from 1 September 2020 through 13 April 2021. This included overlap with the time of the introduction of SARS-CoV-2 vaccines and thus some participants were vaccinated prior to enrollment. The nasal samples were processed at UCSF. Blood samples were processed at Labcorp-Monogram Biosciences, Inc, with plasma stored at  $-80^{\circ}$ C after isolation via centrifugation of heparinized blood. On the day of analysis, the plasma was thawed, centrifuged for 5 minutes at 10 000g, and plated in Quanterix well plates. Assays were performed in accordance with the manufacturer's instructions.

# Measurements

# Blood-Based

We measured SARS-CoV-2 N-antigen in the plasma using the Simoa technology, Quanterix automated paramagnetic microbead-based immunoassay [13]. This technology offers a 1000-fold greater sensitivity than the traditional immunoassay [20, 21]. Quantitative anti–SARS-CoV-2 spike immunoglobulin G (IgG) from the plasma was assessed using a 3-step paramagnetic microbead-based sandwich enzyme-linked immunosorbent assay that also used the Simoa platform. The results were reported as a signal-to-cutoff value, which was determined to be 0.77  $\mu$ g/mL [22]. The lower limit of quantification was 0.098  $\mu$ g/mL, and the upper limit of quantification was 338  $\mu$ g/mL.

### Nasal-Based

We used reverse-transcription polymerase chain reaction (RT-PCR) SARS-CoV-2 RNA and viral culture data from multiple nasal samples over time. Longitudinal RT-PCR RNA and infectious viral shedding data were generated from the selfcollected anterior nasal swabs by the Andino laboratory at UCSF. This lab used the KingFisher platform to target nucleocapsid (N) and envelope (E) gene regions; a SARS-CoV-2 RNA-positive result was defined as having any level of RNA positivity in the N and E gene regions. Cytopathic effect (CPE) was assessed with Vero-hACE2-TMPRSS2 cells in a Biosafety Level 3 laboratory. We defined infection as RT-PCR RNA positivity and defined infectious viral shedding as CPE positive.

#### Questionnaire-Based

We determined symptom status and onset based on data obtained from our interview-administered questionnaires. In addition to symptom data, these questionnaires yielded data on sociodemographic characteristics, medical history, and clinical course of acute coronavirus disease 2019 (COVID-19); the instruments, also used in a related study of COVID-19 recovery, have been described in detail elsewhere [23].

#### **Classification of Infection Status**

We defined true positives as participants with evidence of SARS-CoV-2 infection from at least 1 RT-PCR RNA-positive result from the clinical or research laboratories. We did not include serological data in this assessment because of the potential for false-positive results [24]. We defined true negatives as participants with no longitudinal evidence of a RT-PCR RNA-positive result from the clinical or research laboratories. In our assessment of true negatives, participants were required to have negative IgG antibodies throughout the follow-up period. However, the negative status of vaccinated individuals was not dependent on anti-spike IgG antibodies.

#### **Statistical Analyses**

We assessed the test performance characteristics (sensitivity and specificity) of the SARS-CoV-2 plasma N-antigen against infection (gold standard) and against infectious viral shedding. Evaluation of sensitivity and sensitivity was restricted to the first 14 days of symptom onset, stratified by 0-7 days and 8-14 days. We constructed ROC curves (sensitivity plotted against 1-specificity to varying concentrations of nucleocapsid protein) for each outcome (RNA viral shedding, infectious viral shedding). To demonstrate the range of sensitivity and specificity estimates available from the test, we used the ROC curves to describe values of the N-antigen test that included the manufacturer's cutoff of 1.25 pg/mL, Youden index (sensitivity + specificity - 1) [25], optimized sensitivity, and optimized specificity.We used specimens from the full 28-day observational period to assess the correlation between N-antigen in plasma samples and infectious viral shedding in nasal samples. For analyzing the within-subject correlation between these 2 variables, we used the "analysis of variance" with CPE (infectious viral shedding) as the outcome variable and N-antigen as the predictor variable. The direction of correlation was assessed using a linear regression model. For assessment of between-subject correlation, we used weighted correlation coefficient, using number of observations as weight [26]. All statistical calculations were performed using Stata version 16.1 software (StataCorp LLC, College Station, Texas).

#### RESULTS

#### **Clinical Description of Study Participants**

The analysis cohort contained 80 individuals (Table 1); 44 (55%) were women. Fifty-eight (73%) individuals were true positives, and 22 (27%) individuals were true negatives. Fifty-three of 58 (92%) SARS-CoV-2–infected individuals were symptomatic, and 5 (8%) had asymptomatic infection. Four of 22 (18.2%) individuals without SARS-CoV-2 infection reported some symptoms during the quarantine period even though they did not test positive or seroconvert. A total of

Table 1. Characteristics of Study Participants

|  | SARS-CoV-2 Infection Status           |                                      |                   |
|--|---------------------------------------|--------------------------------------|-------------------|
| Characteristic                           | Infection Present<br>(TP)<br>(n = 58) | Infection Absent<br>(TN)<br>(n = 22) | Total (N = 80)    |
| Classification                           |                                       |                                      |                   |
| Index case                               | 34 (58.6)                             | O (O)                                | 34 (42.5)         |
| Household contact                        | 24 (41.4)                             | 22 (100)                             | 46 (57.5)         |
| Age, y                                   |                                       |                                      |                   |
| Mean (range)                             | 39.92 (16–68)                         | 33.68 (15–56)                        | 37.98 (15–68)     |
| <25 y                                    | 6 (10.3)                              | 6 (27.3)                             | 12 (15.0)         |
| 25–55 y                                  | 42 (72.4)                             | 13 (59.1)                            | 55 (68.7)         |
| 56–65 y                                  | 6 (10.3)                              | 3 (13.6)                             | 9 (11.3)          |
| >65 y                                    | 4 (7.0)                               | 0(0)                                 | 4 (5.0)           |
| BMI, kg/m <sup>2</sup> ,<br>mean (range) | 28.7 (18.4–56.7)                      | 25.24 (18.5–48)                      | 27.8 (18.42–56.7) |
| ≤24.9                                    | 18 (31.0)                             | 13 (59.1)                            | 31 (38.8)         |
| 25-29.9                                  | 19 (32.7)                             | 6 (27.3)                             | 25 (31.2)         |
| ≥30.0                                    | 17 (29.3)                             | 1 (4.5)                              | 18 (22.5)         |
| Not reported/<br>unknown                 | 4 (7.0)                               | 2 (9.1)                              | 6 (7.5)           |
| Biological sex                           |                                       |                                      |                   |
| Female                                   | 32 (55.2)                             | 12 (54.5)                            | 44 (55.0)         |
| Male                                     | 26 (44.8)                             | 10 (45.5)                            | 36 (45.0)         |
| Race/ethnicity                           |                                       |                                      |                   |
| Hispanic/Latino                          | 19 (32.8)                             | 7 (31.8)                             | 26 (32.5)         |
| Non-Hispanic/<br>Latino                  | 39 (67.2)                             | 13 (59.1)                            | 52 (65)           |
| Prefer not to answer                     | 0 (0)                                 | 2 (9.1)                              | 2 (2.5)           |
| Self-reported comorbidities <sup>a</sup> | 18 (31.1)                             | 5 (22.7)                             | 23 (28.7)         |

Data are presented as No. (%) unless otherwise indicated.

Abbreviations: BMI, body mass index; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; TN, true negative; TP, true positive.

<sup>a</sup>Participants self-reported the following comorbidities: autoimmune disease, cancer, diabetes, heart disease, lung disease, and kidney disease.

297 specimens had nasal RT-PCR and corresponding N-antigen results within 24 hours. Of these, 71 (24%) were within 7 days of symptom onset, 110 (37%) were between 8 and 14 days, and the remaining 116 (39%) were  $\geq$ 15 days (Figure 1).

#### Levels and Trends of Plasma N-Antigen

Among true positives, the level of N-antigen in plasma was higher for symptomatic individuals compared to asymptomatic individuals (mean, 37.5 vs 10.8 pg/mL). The mean peak level of N-antigen was 29.6 pg/mL (standard deviation, 108.6), with a subsequent exponential decay over time (Supplementary Figure 1). The median duration of N-antigen positivity in the plasma was 14 days from symptom onset (interquartile range, 8–21 days).

#### Test Characteristics of Plasma N-Antigen Against SARS-CoV-2 Infection

In the evaluation of the test for infection, sensitivity was higher from 0 to 7 days than 8 to 14 days after symptom onset



Figure 1. Description of study cohort. Abbreviation: RT-PCR, reverse-transcription polymerase chain reaction.

(Supplementary Figure 2), whereas specificity remained stable over the 14-day observation period. Using at a cutoff of 1.25 pg/mL from day 0 to day 7, sensitivity was 77.6% (95% confidence interval {CI}, 64%–88.2%), and specificity was 100% (95% CI, 84.6%–100%). From days 0–7 to days 8–14, sensitivity decreased from 77.6% to 43.2% (95% CI, 31.1%–54.5%) whereas specificity was unchanged at 100% (95% CI, 88.1%–100).

While there were points along the ROC curve that optimized both sensitivity and specificity, we found that the following pattern was consistent for further optimizing sensitivity or specificity: decreasing the cutoff increased sensitivity while increasing the cutoff increased specificity (Figure 2). By decreasing the cutoff to 0.48 pg/mL on the ROC curve evaluating infection from day 0 to day 7, for example, sensitivity increased to 86%. We observed a similar pattern from day 8 to day 14: By decreasing the cutoff to 0.26 pg/mL, sensitivity was optimized up to 87.6%. In situations where the cutoff was decreased to optimize or increase sensitivity, there was a concurrent decline in specificity and vice versa.



Figure 2. Receiver operating characteristic curve for N-antigen concentration for detection of severe acute respiratory syndrome coronavirus 2 infection. A total of 71 tests for 0–7 days and 110 tests for 8–14 days after symptom onset were included in the analysis. Abbreviations: ROC, receiver operating characteristic; Sens, sensitivity; Spec, specificity.

# Test Characteristics of Plasma N-Antigen Against Infectious Viral Shedding

### DISCUSSION

In the evaluation of the test for infectious viral shedding, the test had higher sensitivity and lower specificity for infectious viral shedding as compared with infection. Using a cutoff of 1.25 pg/mL to evaluate for infectious viral shedding within 7 days of symptom onset, we found a sensitivity of 100% (95% CI, 88.4%-100%) and specificity of 65% (95% CI, 40.8%-84.6%). By following the ROC curve nearly to its end, we were able to optimize specificity to 100%, but the cutoff increased to 278 pg/mL and sensitivity was very low at 13.3%. Given any cutoff, sensitivity decreased over the 14-day period, which also occurred in the evaluation for infection. Even in the case of infectious viral shedding when we used Youden index, sensitivity still decreased from 96.7% in the first 7 days (0-7 days) to 65% (95% CI, 45.7%-88.1%) in the second 7 days (8-14 days). Returning to the cutoff of 1.25 pg/mL to compare specificity over the 2 periods (0-7 days vs 8-14 days), the estimate was similar at 64%–65% (Figure 3). The test performance for detection of SARS-CoV-2 infectious viral shedding was similar among truly infected individuals rather and the entire study population (Supplementary Figure 3). Over time, we observed that the presence of N-antigen correlated with infectious viral shedding (r = 0.63; P < .01). More details on cutoffs that optimize sensitivity and specificity can be found in Table 2.

In this longitudinally sampled cohort, our findings provide novel evidence in support of the evaluation of N-antigen plasma testing for infectious virus shedding from the nasopharynx. We found that this SARS-CoV-2 N-antigen plasma test was highly sensitive for detecting early infectious viral shedding. The correlation between plasma N-antigen and infectious viral shedding was strong, which was consistent with previous studies describing the first week of acute illness [27-29]. In contrast to RT-PCR testing, antigen testing has the potential to differentiate between the infectious (ie, replication competent) and noninfectious virus [30] [31, 32]. Given the relatively poor specificity to identify infectiousness, this test would not be appropriate for use to test out of quarantine or isolation at the manufacturer's cutoff, but as a highly sensitive test, it may be a viable screening test for community-dwelling individuals to inform their isolation practices.

Compared with other studies evaluating infection, N-antigen plasma testing in our study performed with similar sensitivity and specificity for infection; other studies included outpatients as well as hospitalized individuals and reported sensitivity ranging from 62% to 81.4% and specificity of 93% to 100% [4–10, 33–36]. Several studies assessing performance of nasal N-antigen tests reported higher sensitivity ranging from 80%



Figure 3. Receiver operating characteristic curve for N-antigen concentration for detection of severe acute respiratory syndrome coronavirus 2 infectious viral shedding. A total of 71 tests for 0–7 days and 110 tests for 8–14 days after symptom onset were included in the analysis. Abbreviations: ROC, receiver operating characteristic; Sens, sensitivity; Spec, specificity.

to 98.3% and specificity from 96.8% to 100% [37–40]. In concordance with our findings (specificity of 100% for infection), the outstanding specificity of the N-antigen plasma tests may be ideal in some settings as confirmatory testing, for example point-of-care finger stick testing, pediatric or add-on laboratory testing, and blood banking.

In the evaluation of infection and infectious viral shedding, the ROC curve could be used to identify cutoffs that improve the test performance and optimize it more for screening or confirmatory testing. Depending on the objectives and setting, practitioners can elect how they use the N-antigen plasma test. For example, this test was less sensitive in identifying infected individuals in our nonhospitalized cohort than previously reported with this test in a hospitalized and long-term-care cohort [13]; if a cutoff below current manufacturer recommendations were to be used, however, sensitivity of infection would increase to more acceptable levels in which it could be used as a screening test. Although the specificity of our N-antigen plasma test for infection was outstanding, the specificity for detection of early infectious viral shedding was more moderate, so increasing the cutoff to high levels is a potential solution to optimizing specificity. If specificity were to be outstanding (100%), then this test could be used in hospital-based environments, for example, as an approach to testing out of isolation, instead of alternative imperfect approaches such as cycle threshold values.

This study has several limitations. We included a small number of outpatients, and our findings should be verified in a larger cohort. The study population is not representative of all individuals with SARS-CoV-2, as we studied few asymptomatic individuals as there were few included, and screening asymptomatic individuals would be presumably a useful purpose for this test (eg, blood banks). We did not study variant-specific effects, and the circulating variants during the study period preceded the Delta and Omicron surges in the United States. We also included relatively few vaccinated individuals, and no individuals receiving antiviral therapy; these factors might affect test performance and are becoming more widespread in the current era of SARS-CoV-2.

In conclusion, we demonstrated several ways that the N-antigen plasma test can be used to detect infection and infectious viral shedding. Our evaluation of the test for infection was comparable to other rapid nasal and plasma N-antigen tests available for commercial use. Finally, this study draws on a rigorous sampled longitudinal cohort tested for infectious viral shedding, and the evaluation of N-antigen plasma testing in this cohort provided novel data. Our findings support this test as a suitable screening test for identifying early SARS-CoV-2 infectiousness. Given that the feasibility of this test may make its use preferrable in some settings, this test could be of use in many clinical practice and research settings

 Table 2.
 Summary of N-Antigen Test Performance From Receiver

 Operating Characteristic Curves

| Cutoff                                | Diagnosis<br>of Infection<br>(Within 7 d<br>After<br>Symptom<br>Onset) | Diagnosis<br>of Infection<br>(8–14 d<br>After<br>Symptom<br>Onset) | Diagnosis of<br>Infectious<br>Viral<br>Shedding<br>(Within 7 d<br>After<br>Symptom<br>Onset) | Diagnosis of<br>Infectious<br>Viral<br>Shedding<br>(8–14 d<br>After<br>Symptom<br>Onset) |
|---------------------------------------|--|--|--|--|
| Manufacturer's cutoff                 | 1.3ª   | 1.3  | 1.3  | 1.3  |
| Sensitivity                           | 77.6   | 42.0   | 100.0  | 70.0   |
| Specificity                           | 100.0  | 100.0  | 65.0   | 64.3   |
| Cutoff for<br>highest<br>Youden index | 2.3  | 1.2  | 5.5  | 10.2   |
| Sensitivity                           | 77.6   | 43.2   | 96.7   | 65.0   |
| Specificity                           | 100.0  | 100.0  | 70.0   | 76.2   |
| Cutoff<br>optimizing<br>sensitivity   | 0.5  | 0.3  | 2.3  | 0.3  |
| Sensitivity                           | 85.7   | 87.6   | 100.0  | 90.0   |
| Specificity                           | 72.7   | 34.5   | 65.0   | 14.3   |
| Cutoff<br>optimizing<br>specificity   | 2.3  | 1.2  | 278.4  | 824.0  |
| Sensitivity                           | 77.6   | 43.2   | 13.3   | 5.0  |
| Specificity                           | 100.0  | 100.0  | 100.0  | 97.6   |

Cutoff for N-antigen concentration is shown as picograms per milliliter, and sensitivity and specificity as percentage. Youden index was calculated as (sensitivity + specificity – 1). Sensitivity was calculated as the number of true positives / (number of true positives). Specificity was calculated as the number of true negatives / (number of true negatives + number of false positives).

<sup>a</sup>The manufacturer's cutoff of 1.25 pg/mL is rounded off to 1.3.

if our results are confirmed in larger and more diverse cohorts.

#### Supplementary Data

Supplementary materials are available at *Open Forum Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

#### Notes

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Author contributions. K. A., M. R., J. P.-R., R. D. S., and P. R. recruited participants, coordinated research visits, administered study questionnaires, and collected clinical data. S. L., S. M., S. A. G., J. Y. C., and K. D. performed data entry and quality control and addressed, or adjudicated issues related to data integrity. S. L. maintained the database and oversaw data management. M. G.-K., M. T., and A. Z. performed the RT-PCR test and viral culture in the laboratory. A. C., J. W. W., C. J. P., B. C. Y., and J. L. performed and oversaw the N-antigen assay at Monogram Biosciences and Quanterix laboratory. S. M. and M. C. D. accessed and verified all data, did the data analysis, and drafted the first and final version of the manuscript. J. R. S., D. V. G., G. W. R., S. G. D., M. J. P., R. A. P., J. N. M., and J. D. K. provided in-depth critical review for manuscript revision. All authors reviewed, edited, and approved the manuscript.

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