

# At-home Testing and Risk Factors for Acquisition of SARS-CoV-2 Infection in a Major US Metropolitan Area

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**Background.** Unbiased assessment of the risks associated with acquisition of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is critical to informing mitigation efforts during pandemics. The objective of our study was to understand the risk factors for acquiring coronavirus disease 2019 (COVID-19) in a large prospective cohort of adult residents in a large US metropolitan area.

**Methods.** We designed a fully remote longitudinal cohort study involving monthly at-home SARS-CoV-2 polymerase chain reaction (PCR) and serology self-testing and monthly surveys.

**Results.** Between October 2020 and January 2021, we enrolled 10 289 adults reflective of the Boston metropolitan area census data. At study entry, 567 (5.5%) participants had evidence of current or prior SARS-CoV-2 infection. This increased to 13.4% by June 15, 2021. Compared with Whites, Black non-Hispanic participants had a 2.2-fold greater risk of acquiring COVID-19 (hazard ratio [HR], 2.19; 95% CI, 1.91–2.50;  $P < .001$ ), and Hispanics had a 1.5-fold greater risk (HR, 1.52; 95% CI, 1.32–1.71;  $P < .016$ ). Individuals aged 18–29, those who worked outside the home, and those living with other adults and children were at an increased risk. Individuals in the second and third lowest disadvantaged neighborhood communities were associated with an increased risk of acquiring COVID-19. Individuals with medical risk factors for severe disease were at a decreased risk of SARS-CoV-2 acquisition.

**Conclusions.** These results demonstrate that race/ethnicity and socioeconomic status are the biggest determinants of acquisition of infection. This disparity is significantly underestimated if based on PCR data alone, as noted by the discrepancy in serology vs PCR detection for non-White participants, and points to persistent disparity in access to testing. Medical conditions and advanced age, which increase the risk for severity of SARS-CoV-2 disease, were associated with a lower risk of COVID-19 acquisition, suggesting the importance of behavior modifications. These findings highlight the need for mitigation programs that overcome challenges of structural racism in current and future pandemics.

**Keywords.** COVID-19; SARS-CoV-2; seroprevalence; at-home testing; risk factors.

The severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) pandemic has necessitated frequent decisions regarding prioritization of access to mitigation measures such as testing, contact tracing, housing support, and vaccination among population groups. Considerable disparity in the application of these measures has been observed in the United States, with decreased uptake among younger adults, racial and ethnic minorities, rural populations, individuals with lower

socioeconomic statuses, and certain occupational groups. While several studies have focused on the risk factors for severity of coronavirus disease 2019 (COVID-19) infection, unbiased data are lacking to assess relative population risks regarding the acquisition of COVID-19, which are critical to know to better guide ongoing mitigation efforts by public health authorities, institutions, and health care providers.

While the successful aggregation of SARS-CoV-2 viral polymerase chain reaction (PCR) test results by regional, state, and national public health agencies has permitted comparison of confirmed COVID-19 incidence by geography, age, and racial/ethnic groups, variable access to SARS-CoV-2 testing throughout the pandemic by these same factors, as well as poorer access to testing in many disadvantaged communities, potentially confounds inference related to the risk of different populations acquiring SARS-CoV-2 infection [1, 2]. In addition, given that close to half of all COVID-19 infections may be asymptomatic or only mildly symptomatic, thus not prompting PCR testing, public health agency data based on captured

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viral testing data alone rather than viral and antibody testing may not accurately capture the true incidence. Additionally, the lack of further granularity of testing details, such as employment status, household composition, and medical comorbidities, limits the interpretations of local and national testing trends. These confounding factors have likely led to an under-reporting of SARS-CoV-2 infections. Consequently, unbiased seroprevalence monitoring is very important for obtaining more accurate estimates of infection and transmission as well as determining the risk factors for acquiring COVID-19. Seroprevalence monitoring also contributes to a more refined estimate of the proportion of individuals who have not yet been exposed to SARS-CoV-2 and are not yet vaccinated, and thus constitute the greatest at-risk group of individuals. Cross-sectional seroprevalence studies have documented that 60%–70% of SARS-CoV-2 infections were not clinically detected, but the lack of understanding of the relative timing of infection and potential risks associated with transmission also limits the conclusions of these cross-sectional studies [3, 4].

To inform implementation of risk reduction interventions, we sought to prospectively examine the risks associated with the incidence of SARS-CoV-2 infection in a large cohort of adult residents recruited to be representative of the Boston metropolitan area.

## METHODS

By enrolling a large, generalizable cohort of individuals, representative of the metropolitan Boston area, the aim of this study was to understand the risk factors for acquiring COVID-19 and how confirmed cases of COVID-19 and seroprevalence varied across different socioeconomic statuses, racial/ethnic groups, sexes, age groups, medical comorbidities, and occupations between October 2020 and June 2021. In this monthly cross-sectional study, we tested individuals in the greater metropolitan Boston area for both SARS-CoV-2 PCR and SARS-CoV-2 antibodies. In this article, we present the results, including seroprevalence estimates and cumulative incidence over time based on serologic testing before vaccination and viral testing from adult participants enrolled between October 2020 and January 2021 with longitudinal 6-month follow-up.

### Study Design

A randomly generated cohort of 150 000 adults age 18 years or older who lived within a 45-mile radius of Boston were eligible to enroll in the TestBoston Study. The study was approved by the Mass General Brigham Institutional Review Board. Participants provided informed consent either electronically or via a telephone consent that a qualified study staff member conducted. Consent forms were available in English, Spanish, and Haitian Creole. Participants were offered the opportunity to continue to participate in the longitudinal study with up to

6 months of surveillance testing. The enrolled cohort consisted of individuals normalized to the general population of Massachusetts based on age, gender, and race/ethnicity per state census data [5].

Baseline surveys were completed by enrolled participants followed by monthly questionnaires to collect information on any new COVID-like symptoms, SARS-CoV-2 testing performed outside of the study, and, beginning in January 2021, history of COVID vaccination in the prior 30 days. Self-collection test kits were shipped directly to the participant's home and included an anterior nasal swab for SARS-CoV-2 PCR and a dried blood spot card and lancet for SARS-CoV-2 immunoglobulin G serology. Participants shipped the kit back to the Broad Institute in a pre-addressed, prepaid envelope for processing (instructions in the [Supplementary Data](#)). In addition to monthly surveillance testing irrespective of symptoms, if a study participant developed new symptoms concerning for COVID-19 between scheduled monthly tests or had a known exposure to another individual with COVID-19, the study participant was able to request an additional ad hoc test kit (recruitment methods and laboratory details available in the [Supplementary Data](#)).

### Statistical Analysis

Power analyses set a target enrollment of 10 000 individuals at baseline, followed by 80% of individuals who chose to enroll in the longitudinal portion of the study, assuming a 10% monthly attrition rate during months 3–6. Assuming a baseline seroprevalence of 5%, this sample size was determined to allow us to estimate an overall seroprevalence of COVID-19 infection of 5% or higher (95% CI, 4.5%–5.5%) and to estimate the seroprevalence of COVID-19 infection by age groups, gender, and ethnicity with good precision.

Descriptive statistics (proportions and 95% CIs, means and standard deviations) were performed for all studied variables: demographics, occupation, socioeconomic status as estimated by Area Deprivation Index [6], and comorbidities. Additional analyses including multivariate Cox regression analyses were used to explore factors associated with seroprevalence and viral positivity.

## RESULTS

### Participants, Testing, and Retention

Between October 2020 and January 2021, 102 576 adult residents of metropolitan Boston (estimated population 3 987 800 adults) were invited to join the study, and 10 289 (10%) enrolled and completed at least 1 test kit. The median date of enrollment (interquartile range) was November 23, 2020 (October 20 to December 23, 2020). A total of 7181 (70%) participants were White non-Hispanic, 952 (9.3%) were Hispanic, 925 (9.0%) were Black non-Hispanic, and 889 (8.6%) were

**Table 1. Baseline Characteristics of TestBoston Participants and Metropolitan Boston Population**

Characteristic	TestBoston (n = 10 289), No. (%)	Metro Boston (n = 3 987 800), No. (%)
Sex	...	...
Female	5855 (57)	2 090 300 (52)
Male	4434 (43)	1 897 500 (48)
Age	...	...
18–29 y	1765 (17)	890 600 (22)
30–44 y	3244 (32)	1 036 200 (26)
45–64 y	3534 (34)	1 394 400 (35)
≥65 y	1746 (17)	666 600 (17)
Race and ethnicity	...	...
White	7181 (70)	3 045 800 (76)
Black	925 (9.0)	267 900 (6.7)
Hispanic or Latinx	952 (9.3)	324 100 (8.1)
Asian	889 (8.6)	250 400 (6.3)
Multiple and other races/ethnicities	342 (3.3)	99 600 (2.5)
Neighborhood disadvantage (ADI quintiles)	...	...
0–20th percentile (lowest disadvantage)	4320 (42)	965 400 (24)
20–40th percentile	3052 (30)	983 000 (25)
40–60th percentile	1896 (18)	888 900 (22)
60–80th percentile	799 (7.8)	729 600 (18)
80–100th percentile (highest disadvantage)	222 (2.2)	420 900 (11)
Occupation	...	...
Business, legal, managerial	3252 (32)	1 112 700 (28)
Health care	1526 (15)	285 900 (7.2)
Retail, transportation, and other frontline	745 (7.2)	1 037 500 (26)
Education and sciences	1199 (12)	277 300 (7.0)
Arts, sports, and other	742 (7.2)	170 400 (4.3)
Retired, unemployed, or student	2825 (27)	1 103 800 (28)
Household members	...	...
Living alone	1314 (13)	—
Living with other adults	5645 (55)	—
Living with children	3330 (32)	—
Location of work	...	...
Home or remote	7185 (70)	—
In-person	3104 (30)	—
Medical conditions	...	...
No major comorbidities	5387 (52)	—
Immunocompromise	862 (8.4)	—
Cardiovascular, pulmonary disease, or diabetes	3434 (33)	—
Psychiatric disorder	606 (5.9)	—
Prior COVID-19	...	...
No prior COVID-19	9717 (94)	5 274 300 (97)
PCR-confirmed COVID-19	245 (2.4)	189 800 (3.5)
Serology-detected COVID-19	327 (3.2)	—

Characteristics of TestBoston participants were obtained via self-report and SARS-CoV-2 testing at the time of study entry, with the exception of neighborhood disadvantage, which was determined through geolocation of address to census block group (2010 US Census definitions) and Area Deprivation Index (2018 version). Characteristics for metropolitan Boston were drawn from resident responses to the 2010 US Census (last available at block group level) in the TestBoston study area (45-mile radius around Boston). Cumulative incidence of COVID-19 for towns and cities within the TestBoston study area was obtained from public reporting by the Massachusetts Department of Public Health. Abbreviations: ADI, Area Deprivation Index; COVID-19, coronavirus disease 2019; PCR, polymerase chain reaction.

Asian non-Hispanic. A total of 5855 (57%) participants were female. A total of 1765 (17%) were 18–29 years old, and 1746

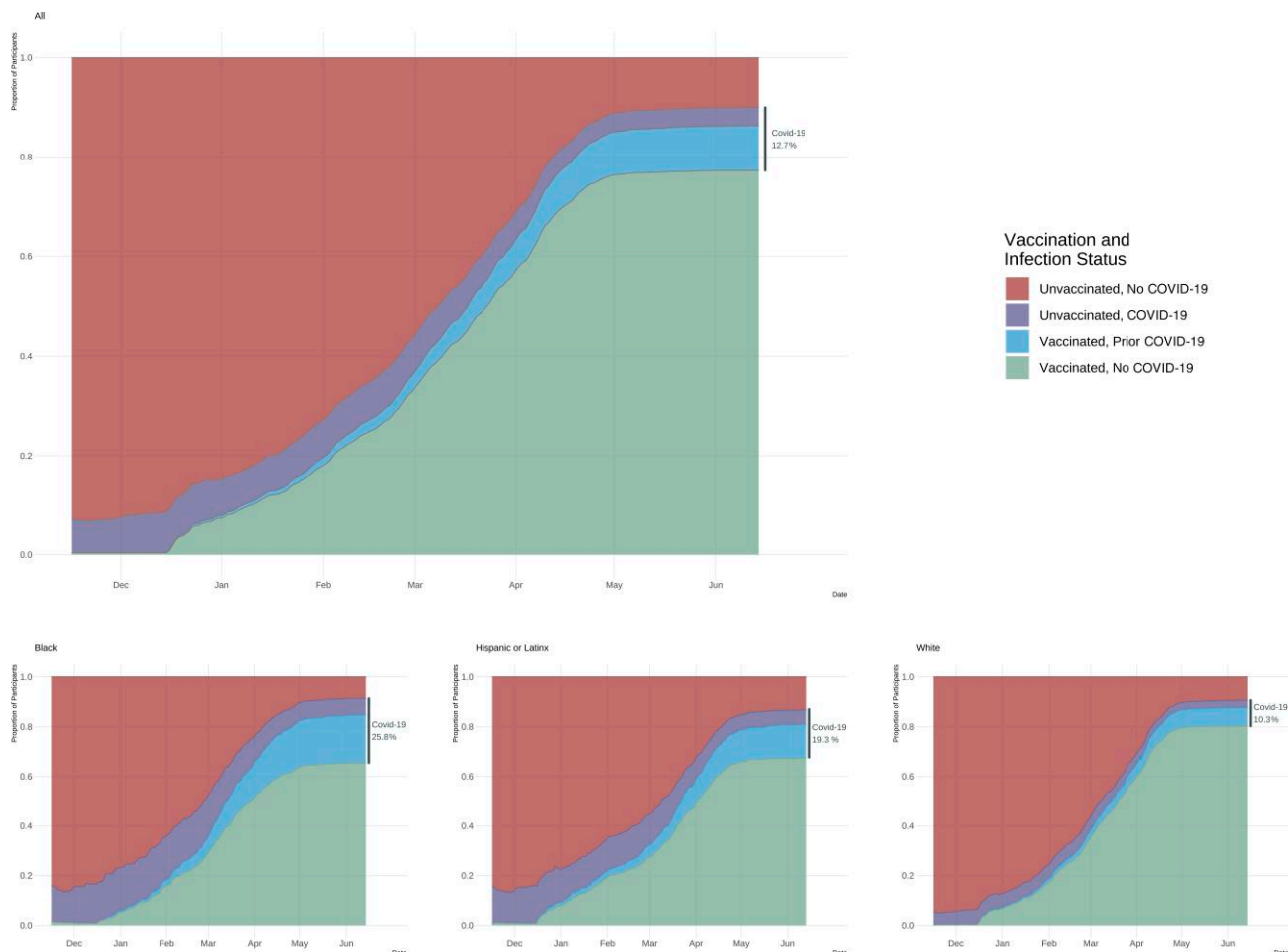
(17%) were age 65 years or older (Table 1). The TestBoston participants had demographic characteristics similar to the metropolitan Boston population, with some observed differences. The TestBoston cohort included more non-White participants (Black [9% in TestBoston vs 6.7% in metropolitan Boston census data]; Hispanic or Latinx [9.3% vs 8.1%]; Asian [8.6% vs 6.3%]; and multiple and other race/ethnicities [3.3% vs 2.5%]), more females (57% vs 52%), more participants in professional occupations (health care [15% vs 7.2%]; education and sciences [12% vs 7%]), and fewer participants in retail, transportation, and other frontline occupations (7.2% vs 26%) than the catchment population. Participants in TestBoston lived in neighborhoods with lower levels of disadvantage and reported being previously diagnosed with COVID-19 at enrollment less frequently than the rate of confirmed COVID-19 in the metropolitan Boston population (2.4% vs 3.5%).

Through the analysis date of June 15, 2021, TestBoston participants were followed for 5174 person-years (median, 6.2 months). Retention for 4 months or more was 56%. During this period, a total of 44 886 viral PCR and 43 628 serology tests were performed by TestBoston, and 15 090 viral PCR tests performed outside of TestBoston were reported by participants. Of the TestBoston tests performed between October 2020 and June 2021, 252 (0.6%) were positive PCR tests and 12 633 (29%) were positive serology tests.

#### SARS-CoV-2 Infection and Vaccination

At the time of study entry, 459 (4.9%) participants had serologic evidence of prior SARS-CoV-2 infection, 2.7 times higher than the 173 who reported having known prior infection. An additional 66 participants were diagnosed by positive PCR at entry. During longitudinal follow-up, an additional 719 (6.9%) infections were detected, with 126 by PCR testing outside of TestBoston, 157 by TestBoston PCR testing, and 613 by serologic testing. The detection of SARS-CoV-2 among TestBoston participants closely followed the epidemic curve for COVID-19 in metropolitan Boston (Supplementary Figure 1), with peak incidence in January 2021.

The cumulative incidence of SARS-CoV-2 infection among TestBoston participants based on serologic testing increased from 5.1% on December 1, 2020, to 12.7% on June 15, 2021. Standardized by age, race/ethnicity, and neighborhood disadvantage, the estimated cumulative incidence among adults in metropolitan Boston increased from 5.1% (95% CI, 4.3%–5.9%) in December 2020 to 12.8% (95% CI, 11.9%–13.7%) in June 2021. In comparison, the cumulative incidence of confirmed COVID-19 in metropolitan Boston, as reported by the Massachusetts Department of Public Health, was 3.1% in December 2020 and 8.4% in June 2021. Non-White TestBoston participants were more likely to have SARS-CoV-2 infection only identified via serology (ie, not clinically recognized or recorded).



**Figure 1.** Cumulative infection and vaccination among TestBoston participants. Date of first positive COVID-19 test (PCR or serology) and first dose of a SARS-CoV-2 vaccine. Participant-reported race and ethnicity at study entry. Abbreviations: COVID-19, coronavirus disease 2019; PCR, polymerase chain reaction; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2.

SARS-CoV-2 vaccination became available to some participants beginning in December 2020, with increasing eligibility through early 2021. A total of 8879 (86.3%) participants received at least 1 vaccine dose during the study period. By June 2021, 9257 (90.0%) of TestBoston participants acquired evidence of immunity to SARS-CoV-2 through vaccination, infection, or both (Figure 1). Standardized to the population, an estimated 89% (95% CI, 88%–91%) of adult residents acquired evidence of immunity to SARS-CoV-2.

#### Risk of Incident SARS-CoV-2 Infection

Among 8532 TestBoston participants without evidence of prior SARS-CoV-2 infection at study entry, 717 (8.4%) acquired COVID-19 during the study period. A multivariate Cox regression analysis (Table 2) showed that individuals aged 30–44 (HR, 0.76; 95% CI, 0.66–0.86;  $P < .001$ ), 45–64 (HR, 0.72; 95% CI, 0.63–0.81;  $P < .001$ ), and 65 years and older (HR, 0.72; 95% CI, 0.61–0.85;  $P < .001$ ) had a significantly lower incidence of SARS-CoV-2 infection compared with 18–29-year-olds.

Conversely, Black non-Hispanic, Asian non-Hispanic, and Hispanic participants had significantly higher incidence (Figure 2). Black non-Hispanics had a 2.2-fold greater risk compared with White non-Hispanics (HR, 2.19; 95% CI, 1.91–2.50;  $P < .001$ ). Hispanics had a 1.5-fold greater risk (HR, 1.52; 95% CI, 1.32–1.71;  $P < .016$ ), and Asian non-Hispanics had a 1.22-fold greater risk (HR, 1.22; 95% CI, 1.04–1.43;  $P < .001$ ) compared with White non-Hispanics. Men had a 16% risk reduction for developing COVID-19 relative to women (HR, 0.84; 95% CI, 0.77–0.92;  $P < .001$ ).

Individuals in the second and third lowest disadvantaged neighborhood communities, as measured by the Area Deprivation Index (ADI) as a marker for socioeconomic status by census block group, were associated with an increased risk of developing COVID-19 (HR, 1.17; 95% CI, 1.04–1.31;  $P = .007$ ; and HR, 1.38; 95% CI, 1.19–1.60;  $P < .001$ ). Living alone was associated with a lower risk of getting COVID-19 as households with adults and children had an increased risk of developing COVID-19 compared with individuals who lived alone

**Table 2. Multivariable Stratified Cox Proportional Hazards Regression Analysis for Incidence of SARS-CoV-2 Infection**

Characteristic	HR	95% CI	P Value
Sex	...	...	
Female	Ref	Ref	
Male	0.84	0.77–0.92	<.001
Age	...	...	
18–29 y	Ref	Ref	
30–44 y	0.76	0.66–0.86	<.001
45–64 y	0.72	0.63–0.81	<.001
≥65 y	0.72	0.61–0.85	<.001
Race and ethnicity	...	...	
White	Ref	Ref	
Black	2.19	1.91–2.50	<.001
Hispanic or Latinx	1.52	1.32–1.75	<.001
Asian	1.22	1.04–1.43	.016
Multiple and other races/ethnicities	1.06	0.82–1.37	.66
Neighborhood disadvantage (ADI quintiles)	...	...	
0–20th percentile (lowest disadvantage)	Ref	Ref	
20–40th percentile	1.01	0.91–1.12	.86
40–60th percentile	1.17	1.04–1.31	.007
60–80th percentile	1.38	1.19–1.60	<.001
80–100th percentile (highest disadvantage)	0.89	0.66–1.20	.44
Medical conditions	...	...	
No major comorbidities	Ref	Ref	
Immunocompromise	0.97	0.83–1.13	.70
Cardiovascular, pulmonary disease, or diabetes	0.91	0.83–1.00	.049
Psychiatric disorder	0.71	0.58–0.88	.002
Location of work	...	...	
Home or remote	Ref	Ref	
In-person	1.32	1.19–1.47	<.001
Occupation	...	...	
Business, legal, managerial	Ref	Ref	
Health care	1.04	0.90–1.21	.58
Retail, transportation, and other frontline	1.02	0.86–1.20	.83
Education and sciences	0.69	0.59–0.80	<.001
Arts, sports, and other	1.08	0.93–1.27	.30
Unemployed	1.06	0.94–1.19	.35
Household members	...	...	
Living alone	Ref	Ref	
Living with other adults	0.92	0.81–1.04	.19
Living with children	1.19	1.04–1.37	.013
Vaccination	...	...	
Unvaccinated	Ref	Ref	
Vaccinated (0–30 d)	0.08	0.05–0.11	<.001
Vaccinated (≥31 d)	0.01	0.00–0.02	<.001

Model was stratified by month of study entry to limit temporal confounding in the context of varying epidemic intensity during the enrollment period. Vaccination status was time-updated, and other characteristics were modeled as fixed as measured at time of study entry.

Abbreviations: ADI, Area Deprivation Index; HR, hazard ratio; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2.

(HR, 1.19; 95% CI, 1.04–1.37;  $P = .013$ ). Individuals who worked outside of the home had an increased risk of acquiring COVID-19 compared with those who worked remotely (HR, 1.32; 95% CI, 1.19–1.47;  $P < .001$ ). Those who worked in education had a lower adjusted risk of developing COVID-19 (HR, 0.69; 95% CI, 0.59–0.80;  $P < .001$ ). Frontline workers

did not have an adjusted increased risk (HR, 1.02; 95% CI, 0.86–1.20;  $P = .83$ ). Vaccination showed a 97.2% rate of effectiveness during this study period (HR, 0.01; 95% CI, 0.00–0.02;  $P < .001$ ).

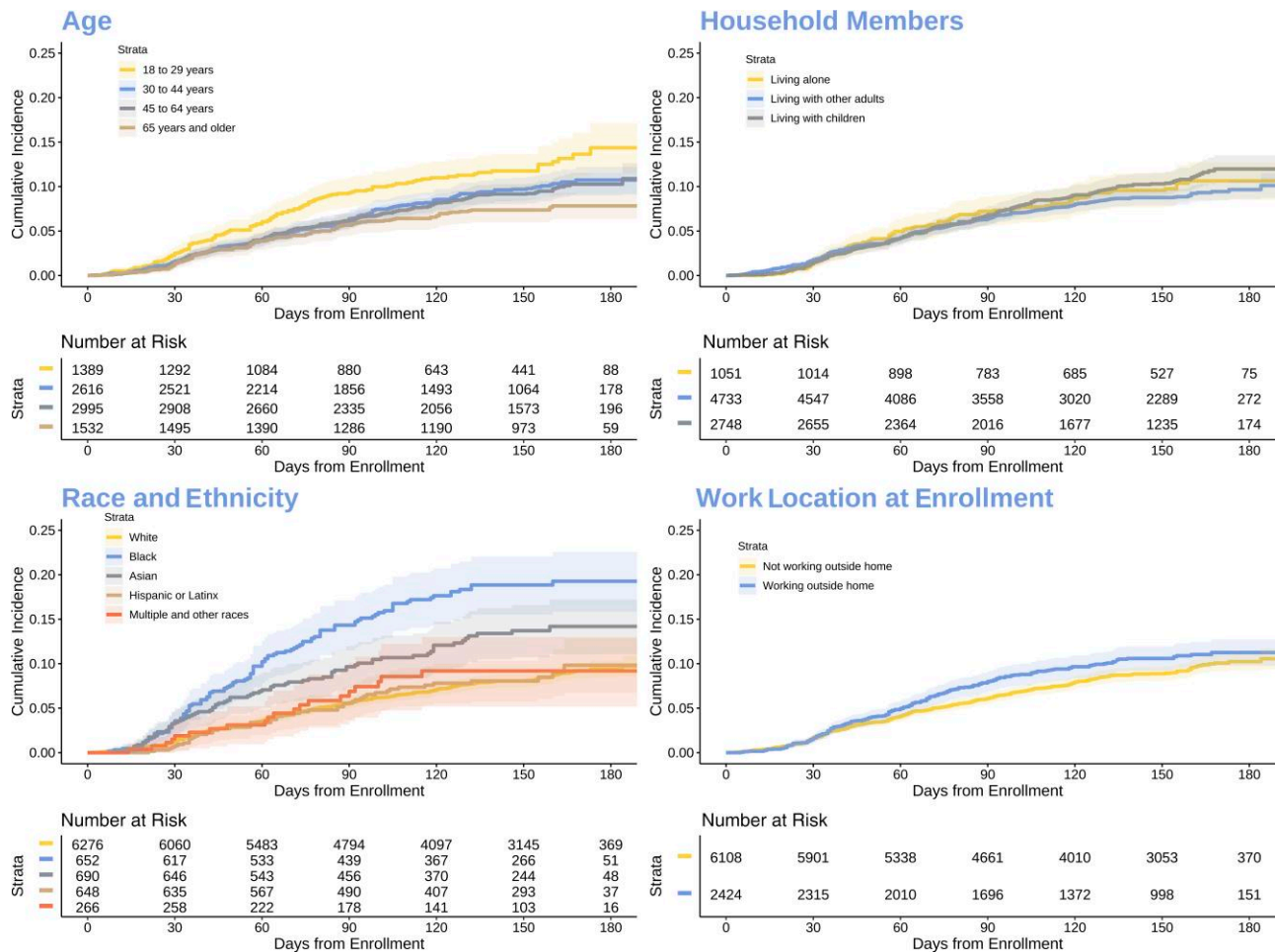
In contrast to medical risk factors for increased severity of infection, individuals with major medical comorbidities did not have an increased risk of acquiring SARS-CoV-2. In fact, individuals with cardiovascular disease, pulmonary disease, obesity, or diabetes had a lower adjusted risk of acquiring COVID-19 (HR, 0.91; 95% CI, 0.83–1.00;  $P = .049$ ), as did individuals with underlying psychiatric disorders such as depression and anxiety (HR, 0.71; 95% CI, 0.58–0.88;  $P = .002$ ).

## DISCUSSION

TestBoston is one of the largest longitudinal, remote, at-home direct-to-patient COVID-19 testing studies in the United States, established to follow a cohort of >10 000 individuals who were representative of the residents of Boston, a US metropolitan city with a diverse population, in order to assess the risk factors associated with development of SARS-CoV-2 infection. There are significant impediments to participating in research studies that disproportionately impact the most vulnerable individuals, particularly those of Black race and Hispanic/Latinx ethnicity, due to language barriers, mistrust in medical research, and time and travel commitments that cannot be accommodated by shift workers, in particular, and those of lowest socioeconomic status. Therefore, this study was designed as a fully remote study that was equitable, scalable, and allowed for a wide geographic catchment area that enabled recruitment and retention of those individuals who are often not represented in these research studies.

As of June 15, 2021, 90% of individuals living in the greater Boston area had previously been infected by SARS-CoV-2 and/or were vaccinated. While reinfection can occur despite prior infection and vaccine breakthrough infections occur, those 90% of individuals are much less likely to be at risk of severe COVID-19 if they subsequently acquire SARS-CoV-2 infection [7–11]. Among the remaining individuals who are therefore at higher risk for severe COVID-19 infection, Black and/or Hispanic individuals are disproportionately represented. This disparity is due to considerably lower vaccination rates in Black and Hispanic individuals, as their rates of prior COVID-19 infection were higher than those of other racial and ethnic groups.

Black race, Asian race, and Hispanic ethnicity were strongly associated with seropositivity and COVID-19 cases, echoing broader racial inequities in testing and consistent with previous reports on the disproportionate burden of disease among some racial and ethnic subgroups seen in the United States during the COVID-19 pandemic [12–14]. Here, we now confirm that this disparity is significantly underestimated if based on



**Figure 2.** Comparison of cumulative incidence of SARS-CoV-2 infection among TestBoston participants without prior infection at study entry. Characteristics as reported by participants at study entry. Abbreviation: SARS-CoV-2, severe acute respiratory syndrome coronavirus 2.

SARS-CoV-2 PCR testing data alone, likely due to unequal access to testing. While there was a 2-fold greater positivity rate based on serology compared with PCR testing for White non-Hispanic and Hispanic individuals, there was a 3.5-fold and 4-fold difference for Black and Asian non-Hispanic individuals.

Twenty-six percent of Black non-Hispanics were seropositive for SARS-CoV-2 before vaccination, but only 7% had tested positive at any time via viral PCR testing either during the study or before study enrollment. Conversely, only 9% of White non-Hispanic individuals were SARS-CoV-2 seropositive before vaccination, and nearly half of those cases had been detected via viral PCR testing, suggesting greater access to testing for White non-Hispanic individuals in the greater Boston area. This disparity in testing is consistent with what has been seen throughout other metropolitan cities throughout the United States, showing the marginalization of majority Black areas due to economic disenfranchisement and limited health-promoting attributes [1, 15–17]. Notably, in New York

City, efforts to increase testing availability in order to decrease the disparity in the placement of testing sites has begun to make an impact [17]. It is imperative that access to testing be improved for vulnerable groups and that racial and ethnic disparities in access to testing be decreased. This is critical for identifying cases and preventing the spread of COVID-19 among these groups. TestBoston has demonstrated that at-home testing is one way to decrease barriers and expand testing [18].

Disparities in testing result in disparities in recognizing cases of COVID-19. The early identification of cases when an individual is asymptomatic or only mildly symptomatic is critical in preventing severe COVID-19 infection and associated morbidities and mortality now that treatment options such as monoclonal antibodies are available to decrease an individual's risk of severe COVID-19 and hospitalization. These study results highlight the importance of understanding the risk factors associated with the acquisition of SARS-CoV-2 infection and the associated negative impact that testing inequities has on

individuals and certain communities receiving the needed medical interventions to prevent ongoing and more severe disease.

We know in theory that an individual's medical comorbidities and behaviors can affect the risk of acquisition of COVID-19, but the data to date have focused more on the risk factors for severity of disease rather than on the risk factors for acquisition. Risk factors of acquisition are critical to elucidating who we should target for mitigation efforts as well as on whom we should focus the emerging outpatient treatment interventions, including monoclonal antibodies and oral antivirals, as they become available, to attenuate the severity of disease, limit ongoing transmission to those individuals who are at higher risk for hospitalization, and minimize long COVID [19].

Younger individuals (18–29 years old) had a higher adjusted risk of acquiring COVID-19 compared with individuals who were age 30 years or older, which may have reflected social behaviors during this period of the pandemic. However, the older age groups had a lower proportion of COVID-19 cases detected by PCR testing and a greater proportion of cases detected only by antibody testing before vaccination. This highlights the greater access to viral testing programs at colleges and universities and certain companies or places of work. This has direct negative clinical impacts as those who are older and have a higher risk of severity of disease were less likely to have had viral COVID-19 testing, which significantly disadvantages them in terms of potential outpatient treatment options to prevent hospitalizations.

Surprisingly, while cardiovascular and pulmonary comorbidities including obesity, diabetes, asthma, and chronic obstructive pulmonary disease, as well as immunosuppressive conditions such as hematologic malignancy, solid organ transplant, or being on immunosuppressive medications, increase an individual's risk for severe illness from SARS-CoV-2 infection, this study found that these medical conditions do not increase one's risk of acquiring COVID-19. Having a psychiatric disorder was seen to be associated with lower risk of acquiring COVID-19, in contrast to severe disease, suggesting that the risk of acquisition can be mitigated by behavior.

The strengths of this study include the frequent, repeat paired SARS-CoV-2 viral and antibody testing of a wide range of individuals from differing racial and ethnic groups, ages, socioeconomic statuses, occupations, and household compositions that are representative of a greater metropolitan city, Boston. This granularity of detail on such a large representative cohort sets this study apart and expounds upon the more limited data that are monitored at the city, state, and national levels. Our study design avoided the biases that have been seen in many of the other large COVID-19 studies because it was not limited to participants who were only testing due to being symptomatic, having risk factors for developing severe COVID, or presenting with severe disease. We were mindful

that data quality matters as much as, if not more than, data quantity and ensured that our findings were not a victim of the "big data paradox," a mathematical tendency of big data sets to minimize errors due to small sample size while magnifying errors linked to systematic biases that make the sample a poor representation of the larger population [20, 21]. As a result of our study design capturing the demographics and composition of a metropolitan US city, we were able to learn key findings about the risks of acquisition of SARS-CoV-2 infection at an individual level as well as a community level. The limitations of this study include the fact that the results analyzed for this study stop in mid-June 2021, before the Delta and now Omicron variants became the predominant circulating strains in this geographic area. Other limitations include that asymptomatic viral PCR testing was only offered once monthly and the inability to detect new cases via serology testing postvaccination.

The COVID-19 pandemic has further highlighted the social and racial inequities that exist in health care and has brought these discussions to the forefront of public health. It has highlighted that health equity is still not a reality as COVID-19 has unequally affected many racial and ethnic minority groups. Testing only captured a fraction of the cases, which resulted in an immense inequity of counting of cases in certain populations. As we face SARS-CoV-2 becoming an endemic viral infection, it is paramount to continue understanding the risk factors for acquisition rather than just severity of illness and which of those factors can be mitigated, as preventing acquisition of infection is the best intervention to prevent severe illness. These data also demonstrate that a major determinant of risk of acquisition is behavior and that a major risk-reducing intervention is behavior modification. For example, in younger individuals, behavior dominates their risk for acquisition, while in individuals with significant medical comorbidities, any possibility of increased acquisition may be offset by active behavioral modifications to minimize exposure. Thus, the importance of behavioral modification during a pandemic cannot be overemphasized, in contrast to medical comorbidities and older age, which cannot be modified.

Taken together, these results suggest both that the consequences of structural racism on risk must be considered during the development of mitigation programs for current and future pandemics and that an important component of these mitigation programs must be to educate and encourage risk-modifying behaviors. Given that Black and Hispanic frontline workers are overrepresented in positions associated with higher risk for COVID-19, proposed behavior modifications, while critical, should also consider such barriers to risk mitigation in the workplace [22]. While the actual risk factors may change as the status of the pandemic continues to evolve, including waning immunity from vaccination, the rise of variants, the availability of testing, and the changing behavioral norms of

local communities and nations, these principles will likely remain firm for the foreseeable future.

### 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.

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**Author contributions.** A.W., D.H., and L.C. designed the study. A.W., A.K., S.N.M., C.K., H.L., D.H., and L.C. conducted the study, enrolled participants, and collected the data. H.L., J.L., P.M., S.G., and D.H. processed the lab samples (either the PCR respiratory samples or dried blood spot serology samples). A.W., S.D.P., J.B., J.L., P.M., M.O., J.G., N.S., D.H., and L.C. provided scientific interpretation of the data. A.W. and S.D.P. performed the statistical analysis. A.W., S.D.P., and L.C. had full access to all the data in the study and take responsibility for the integrity of the data and the accuracy of the data analysis. A.W., S.D.P., D.H., and L.C. drafted the manuscript. All authors revised the manuscript critically for important intellectual content and approved the final version of the manuscript.

**Patient consent.** The authors attest that they are in compliance with the ethical standards of the Helsinki Declaration and human studies committees of the authors' institutions. Informed consent was obtained from all patients or their authorized representatives.

### References

1. Dryden-Peterson S, Velásquez GE, Stopka TJ, Davey S, Lockman S, Ojikutu BO. Disparities in SARS-CoV-2 testing in Massachusetts during the COVID-19 pandemic. *JAMA Netw Open* **2021**; 4:e2037067.
2. Bigelow BF, Saxton RE, Flores-Miller A, et al. Community testing and SARS-CoV-2 rates for Latinxs in Baltimore. *Am J Prev Med* **2021**; 60:e281–6.
3. Rogawski McQuade ET, Guertin KA, Becker L, et al. Assessment of seroprevalence of SARS-CoV-2 and risk factors associated with COVID-19 infection among outpatients in Virginia. *JAMA Netw Open* **2021**; 4:e2035234.
4. Roederer T, Mollo B, Vincent C, et al. Seroprevalence and risk factors of exposure to COVID-19 in homeless people in Paris, France: a cross-sectional study. *Lancet Public Heal* **2021**; 6:e202–9.
5. US Census Bureau. 2010 census of population and housing. Available at: <https://www.census.gov/prod/cen2010/cph-2-1.pdf>. Accessed January 31, 2022.
6. Kind 2018, University of Wisconsin School of Medicine Public Health. 2015 Area Deprivation Index v2.0. Available at: <https://www.neighborhoodatlas.medicine.wisc.edu/>. Accessed June 23, 2021.
7. Abu-Raddad LJ, Chemaitelly H, Ayoub HH, et al. Association of prior SARS-CoV-2 infection with risk of breakthrough infection following mRNA vaccination in Qatar. *JAMA* **2021**; 326:1930–9.
8. Letizia AG, Ge Y, Vangeti S, et al. SARS-CoV-2 seropositivity and subsequent infection risk in healthy young adults: a prospective cohort study. *Lancet Respir Med* **2021**; 9:712–20.
9. Baden LR, El Sahly HM, Essink B, et al. Efficacy and safety of the mRNA-1273 SARS-CoV-2 vaccine. *N Engl J Med* **2021**; 384:403–16.
10. North CM, Barczak A, Goldstein RH, et al. Determining the incidence of asymptomatic SARS-CoV-2 among early recipients of COVID-19 vaccines (DISCOVER-COVID-19): a prospective cohort study of healthcare workers before, during and after vaccination. *Clin Infect Dis* **2022**; 74:1275–8.
11. Brown CM, Vostok J, Johnson H, et al. Outbreak of SARS-CoV-2 infections, including COVID-19 vaccine breakthrough infections, associated with large public gatherings — Barnstable County, Massachusetts, July 2021. *MMWR Morb Mortal Wkly Rep* **2021**; 70:1059–62.
12. Magesh S, John D, Li WT, et al. Disparities in COVID-19 outcomes by race, ethnicity, and socioeconomic status: a systematic-review and meta-analysis. *JAMA Network Open* **2021**; 4:e2134147.
13. Mackey K, Ayers CK, Kondo KK, et al. Racial and ethnic disparities in COVID-19-related infections, hospitalizations, and deaths. *Ann Intern Med* **2021**; 174:362–73.
14. Pan AP, Khan O, Meeks JR, et al. Disparities in COVID-19 hospitalizations and mortality among Black and Hispanic patients: cross-sectional analysis from the greater Houston metropolitan area. *BMC Public Health* **2021**; 21:1330.
15. Poteat T, Millett GA, Nelson LRE, Beyrer C. Understanding COVID-19 risks and vulnerabilities among Black communities in America: the lethal force of syndemics. *Ann Epidemiol* **2020**; 47:1–3.
16. Williams DR, Collins C. Racial residential segregation: a fundamental cause of racial disparities in health. *Public Health Rep* **2001**; 116:404–16.
17. Grigsby-Toussaint DS, Shin JC, Jones A. Disparities in the distribution of COVID-19 testing sites in Black and Latino areas in New York City. *Prev Med* **2021**; 147:106463.
18. Naz-McLean S, Kim A, Zimmer A, et al. Feasibility and lessons learned on remote trial implementation from TestBoston, a fully remote, longitudinal, large-scale COVID-19 surveillance study. *PLoS One* **2022**; 17:e0269127.
19. Bierle DM, Ganesh R, Wilker CG, et al. Influence of social and cultural factors on the decision to consent for monoclonal antibody treatment among high-risk patients with mild-moderate COVID-19. *J Prim Care Community Health* **2021**; 12:21501327211019280.
20. Meng XL. Statistical paradises and paradoxes in big data (I): law of large populations, big data paradox, and the 2016 US presidential election. *Ann Appl Stat* **2018**; 12:685–726.
21. Bradley VC, Kuriwaki S, Isakov M, Sejdinovic D, Meng X-L, Flaxman S. Unrepresentative big surveys significantly overestimated US vaccine uptake. *Nature* **2021**; 600:695–700.
22. Goldman N, Pebley AR, Lee K, Andrasfay T, Pratt B. Racial and ethnic differentials in COVID-19-related job exposures by occupational standing in the US. *PLoS One* **2021**; 16:e0256085.