1	SARS-CoV-2 incidence and risk factors in a national, community-based prospective
2	cohort of U.S. adults
3	Denis Nash ^{1,2,*} , Madhura S. Rane ¹ , McKaylee M. Robertson ¹ , Mindy Chang ¹ , Sarah Gorrell
4	Kulkarni ¹ , Rebecca Zimba ^{1,2} , William You ¹ , Amanda Berry ¹ , Chloe Mirzayi ^{1,2} , Shivani
5	Kochhar ¹ , Andrew Maroko ^{1,3} , Drew A. Westmoreland ¹ , Angela M. Parcesepe ^{1,4,5} , Levi
6	Waldron ^{1,2} , and Christian Grov ^{1,6}
7	¹ Institute for Implementation Science in Population Health (ISPH), City University of New
8	York (CUNY); New York City, New York USA
9	² Department of Epidemiology and Biostatistics, Graduate School of Public Health and Health
10	Policy, City University of New York (CUNY); New York City, New York USA
11	³ Department of Environmental, Occupational, and Geospatial Health Sciences, Graduate
12	School of Public Health and Health Policy, City University of New York (CUNY); New York
13	City, New York USA
14	⁴ Department of Maternal and Child Health, Gillings School of Public Health, University of
15	North Carolina, Chapel Hill, NC, USA
16	⁵ Carolina Population Center, University of North Carolina at Chapel Hill, Chapel Hill, NC,
17	USA
18	⁶ Department of Community Health and Social Sciences, Graduate School of Public Health
19	and Health Policy, City University of New York (CUNY); New York City, New York USA
20	*CORRESPONDING AUTHOR:
21	Denis Nash, Ph.D., MPH
22	CUNY Graduate School of Public Health and Health Policy
23	55 W. 125th St., 6th Floor
24	New York, NY USA 10027
25	Email: <u>denis.nash@sph.cuny.edu</u>
26	Running title: SARS-CoV-2 incidence in a U.S. cohort

© The Author(s) 2022. Published by Oxford University Press on behalf of the Infectious Diseases Society of America. All rights reserved. For permissions, please e-mail: journals.permissions@oup.com This article is published and distributed under the terms of the Oxford University Press, Standard Journals Publication Model (https://academic.oup.com/journals/pages/open_access/funder_policies/chorus/standard_publication_ 1 model)

1 ABSTRACT

2 Background: Prospective cohort studies of SARS-CoV-2 incidence complement case-

3 based surveillance and cross-sectional seroprevalence surveys.

4 **Methods**: We estimated the incidence of SARS-CoV-2 infection in a national cohort of 6,738

5 U.S. adults, enrolled March-August 2020. Using Poisson models, we examined the

6 association of social distancing and a composite epidemiologic risk score with

7 seroconversion. The risk score was created using LASSO regression to identify factors

8 predictive of seroconversion. The selected factors were household crowding, confirmed case

9 in household, indoor dining, gathering with groups \geq 10, and no masking in gyms/salons.

10 **Results:** Among 4,510 individuals with ≥1 serologic test, 323 (7.3%, 95% confidence interval

11 [CI] 6.5%-8.1%) seroconverted by January 2021. Among 3,422 participants seronegative in

12 May-September 2020 and retested during November 2020-January 2021, 161

13 seroconverted over 1,646 person-years of follow-up (9.8 per 100 person-years [95%CI 8.3-

14 11.5]). Seroincidence rate was lower among females compared to males (IRR: 0.69, 95% CI

15 0.50-0.94) and higher among Hispanic (IRR: 2.09, 95% CI 1.41-3.05) participants compared

16 to White non-Hispanic. In adjusted models, participants who reported social distancing with

people they did not know (IRR_{always vs. never}: 0.42, 95% CI 0.20-1.0) and with people they knew

18 (IRR_{alwavs vs. never} 0.64, 95%CI 0.39-1.06; IRR_{sometimes vs. never} 0.60, 95% CI 0.38-0.96) had lower

19 seroconversion risk. Seroconversion risk increased with epidemiologic risk score (IRR_{medium}

20 vs. low 1.68, 95% CI 1.03-2.81; IRR_{high vs. low} 3.49, 95% CI 2.26-5.58). Only 29% of those who

21 seroconverted reported isolating and 19% were asked about contacts.

Conclusion: Modifiable risk factors and poor reach of public health strategies drove SARS CoV-2 transmission across the U.S.

KEYWORDS: COVID-19; serology, seroconversion, asymptomatic infection, physical
 distancing; natural history study, epidemiologic study, essential workers, public health

26 interventions, community transmission

1 INTRODUCTION

2 A major challenge of controlling community transmission of SARS-CoV-2 is the 3 asymptomatic and pre-symptomatic spread of infection [1,2], including by fully-vaccinated individuals [3]. One national study in the United States (U.S.) prior to the vaccine era 4 5 estimated five undiagnosed infections for every diagnosed case [2]. While SARS-CoV-2 is 6 transmitted from person-to-person via airborne and droplet spread, to date, the incidence of 7 SARS-CoV-2 infection and its risk factors have not been adequately characterized by routine case-based surveillance or by cross-sectional seroprevalence studies [4-6]. It is critical for 8 prospective studies to investigate COVID-19's evolving epidemiology, risk factors for SARS-9 CoV-2 incidence in communities, uptake and impact of non-pharmaceutical interventions 10 11 (NPIs) [7], and the reach of public health strategies aimed at controlling community 12 transmission, including testing, guarantine, isolation, contact tracing, and vaccination. Globally, few community-based prospective epidemiologic studies of SARS-CoV-2 incidence 13 and risk factors have been undertaken. One systematic review found 18 prospective 14 observational studies of SARS-CoV-2 that employed serologic or polymerase chain reaction 15 (PCR) testing [8]. Most studies focused on healthcare workers or other occupational groups 16 17 and individuals in congregate settings [8]. While national community-based prospective cohort studies have been conducted in the United Kingdom [9-11], such studies have been 18 scarce in the U.S. Community-based studies with longitudinally collected biomarkers data 19 20 can help inform implementation of public health responses and policies, both for the current pandemic and future ones. 21 In March 2020, we launched the prospective Communities, Households and SARS-CoV-2 22

Epidemiology (CHASING) COVID Cohort [12]. We describe the incidence of SARS-CoV-2
infection and risk factors for SARS-CoV-2 seroconversion during May 2020-January 2021,
and the reach and uptake of public health strategies aimed at controlling community spread
among those who seroconverted.

- 27
- 28

1 METHODS

2 Recruitment

- 3 We used internet-based strategies [13,14] to recruit a geographically and socio-
- 4 demographically diverse cohort of adults into longitudinal follow-up with at-home specimen
- 5 collection. Informed consent was obtained from all participants prior to enrollment. To be
- 6 eligible for inclusion in the cohort, individuals had to: 1) reside in the U.S. or a U.S. territory;
- 7 2) be \geq 18 years old; 3) provide a valid email address; and 4) demonstrate early engagement
- 8 in study activities (provide baseline specimen or complete of >1 recruitment/enrollment visit).
- 9 Details of the study design and recruitment procedures are described elsewhere [15]. The
- 10 full cohort includes participants from all 50 U.S. states, the District of Columbia, Puerto Rico,
- and Guam (Supplemental Materials 1, Figure S2). Of the 6,738 participants in the full cohort,
- 12 4,510 (67%) had at least one serologic test and comprised the study population for this
- 13 analysis (Supplemental Materials 1, Table S1).

14 Data collection

15 Cohort recruitment and multiple rounds of interviews occurred between March 28-August 21,

16 2020. Demographic and COVID-19 related risk factors were collected at baseline. From

17 three follow-up interviews between August-November 2020, we obtained repeated

18 measurements of COVID-19 symptoms, laboratory testing (PCR or serologic),

19 hospitalizations, use of NPIs such as mask use and social distancing, and public health

20 strategies such as quarantine, isolation, and contact tracing.

21 During May-September 2020 (Period 1) and November 2020-January 2021 (Period 2),

22 participants were invited to complete serologic testing using an at-home dried blood spot

- 23 (DBS) specimen self-collection kit. DBS cards were returned to the study laboratory
- 24 (Molecular Testing Laboratories [MTL], Vancouver, Washington) via the U.S. Postal Service
- 25 using a self-addressed, stamped envelope containing a biohazard bag™. All DBS
- specimens were tested by the study laboratory for total antibodies (Total Ab) using the Bio-
- 27 Rad Platelia test for IgA, IgM, and IgG which targets the SARS-CoV-2 nucleocapsid protein
- 28 (manufacturer sensitivity 98.0%, specificity 99.3%) [16]. Other studies have independently

1 validated this assay and found average sensitivity and specificity of 91.7% and 98.8%,

2 respectively [17–19]. This assay was also validated for use with DBS by the study

3 laboratory, which found 100% sensitivity and 100% specificity (Supplemental Materials 2).

4 Outcomes

5 *Cumulative incidence of SARS-CoV-2 infection.* Among participants who underwent 6 serologic testing, we estimated the serology-based cumulative incidence of SARS-CoV-2 as 7 the number of individuals with a positive Total Ab test in either of the two time periods 8 divided by the total number of persons with one or more Total Ab tests. We adjusted 9 cumulative incidence estimates for laboratory test error, assuming a sensitivity of 91.7% and 10 a specificity of 98.8% [17–19] using the following formula [20]:

 $Adjusted \ cumulative \ incidence = \frac{Crude \ cumulative \ incidence + Specificity - 1}{Sensitivity + Specificity - 1}$

Observed SARS-CoV-2 seroconversion. Among individuals with two Total Ab tests, an 11 observed seroconversion was defined as a negative Total Ab test in Period 1 followed by a 12 positive Total Ab test in Period 2. We estimated person-years of follow-up using the 13 specimen collection dates in Periods 1 and 2. We used the date the laboratory received the 14 15 sample for missing collection dates. The seroconversion date was assigned as the midpoint between the first and second specimen collection dates for person-time calculations. To 16 address the possibility of misclassifying some individuals with recent infections who had not 17 vet seroconverted by the time of the second antibody test, we conducted a sensitivity 18 19 analysis including reports of recent positive SARS-CoV-2 PCR or rapid test in the incidence 20 estimate.

21 Exposures

Individual-level COVID-19 risk factors. Individually and as part of a composite score, we
 considered epidemiologic risk factors for SARS-CoV-2 reported by participants prior to
 specimen collection, including: Household factors (household crowding defined as ≥4 people
 living in a multi-family dwelling, having a child in the household, and having a confirmed
 COVID-19 case in a household member prior to participant testing positive); spending time

1 in public places (attending mass gatherings, indoor dining in a restaurant/bar, outdoor dining 2 at a restaurant/bar, visiting places of worship, or visiting public parks/pools); mask use 3 indoors (for grocery shopping, visiting non-household members, at work, and in 4 salons/gyms); mask use outdoors; gathering in groups with 10 people; travel during the 5 pandemic (air travel and public transit use); and individual-level factors that may increase the 6 risk of severe COVID-19 (substance use, binge drinking, and comorbidities). 7 Global social distancing assessment. While social distancing in specific scenarios is 8 addressed in some of the above individual risk factors, we were interested in the association between social distancing in general and incident SARS-CoV-2 infection. We asked two 9 10 global guestions on social distancing: "In the past month, how often have you practiced 11 social distancing with: a) people you know and b) people you do not know," with possible response options of Always, Sometimes, or Never. These assessments were not included in 12 13 the calculation of the composite risk score. Composite score of COVID-19 risk factors. We computed a composite COVID-19 risk score 14

as many of the above COVID-19 risk factors may be highly correlated. We applied Least 15 Absolute Shrinkage Selection Operator (LASSO) regression to select the set of risk factors 16 17 which best predicted seroconversion [21]. The LASSO model selected household crowding, having a confirmed COVID-19 case in a household member, indoor dining in a 18 bar/restaurant, gathering with groups of ≥ 10 , and no mask use indoors in salons/gyms as the 19 most predictive of seroconversion in our cohort. Scores were assigned to each participant 20 based on their engagement in the risk factors selected by the LASSO model and were 21 normalized between 0 and 100. High scores indicate engagement in high-risk activities 22 (Details in Supplemental Materials 1, Statistical Appendix). The composite score was divided 23 24 into tertiles for analysis.

25 Statistical analysis

Cumulative incidence estimates were stratified by baseline characteristics and epidemiologic
 risk factors. Crude and adjusted Incidence Rate Ratios (IRRs) of seroincidence and
 associated 95% confidence intervals (CIs) were estimated using Poisson regression. We

- 1 examined crude seroconversion rates by sociodemographic characteristics and each risk
- 2 factor. Finally, we separately modeled three exposure variables: 1) social distancing with
- 3 "people you know" (always/sometimes/never); 2) social distancing with "people you don't
- 4 know" (always/sometimes/never); and 3) the composite COVID-19 risk score
- 5 (high/medium/low). Two multivariable models were constructed for each exposure variable,
- 6 adjusting for age, gender, race/ethnicity and comorbidities (Model 1); and further controlling
- 7 for changes in community-level COVID-19 transmission (Model 2). All data were cleaned
- 8 and analyzed in R (version 4.0.3) and SAS (V9.4).

9 Ethical Approval

- 10 The study protocol was approved by the Institutional Review Board at the City University of
- 11 New York (CUNY) and all methods were performed in accordance with relevant guidelines
- 12 and regulations.
- 13 **RESULTS**

14 Sample characteristics

- 15 Of the 4,510 participants who tested at least once, 3,605 (80%) tested at both time points
- 16 (<u>Table 1</u>). A total of 4,232 persons underwent serologic testing in Period 1, and 3,883 in
- 17 Period 2 (Supplemental Materials 1, Table S1). Differences between participants testing in
- 18 Period 1 and Period 2 were negligible (Supplemental Materials 1, Table S1). The median
- 19 time between specimen collection dates for participants providing specimens for both
- serologic tests was 190 days (IQR 152-201) (Supplemental Materials 1, Figure S1).

21 Cumulative incidence of SARS-CoV-2 as of January 31, 2021

- 22 We observed 323 unique seropositives among the 4,510 participants who tested at least
- 23 once during follow-up, for overall crude and adjusted serology-based cumulative incidence
- 24 estimates of 7.3% (95% CI 6.5%-8.1%) and 6.7% (95% CI 5.9%-7.6%), respectively
- 25 (Supplemental Materials 1, Table S2).

26 SARS-CoV-2 seroincidence, May 2020-January 2021

- 27 There were 3,422 seronegative participants in Period 1 with a subsequent serologic test in
- 28 Period 2. There were 161 observed seroconversions over 1,646 person years of follow-up,

for an overall incidence rate of 9.8 per 100 person-years (95% CI 8.3-11.5) (Table 2). The
rate of incident SARS-CoV-2 infection was lower for females compared to males (IRR=0.69,
95% CI 0.50-0.94), and higher for Hispanic (IRR=2.09, 95% CI 1.41-3.05) and Black nonHispanic (IRR=1.69, 95% CI 0.96-2.82) compared with White non-Hispanic participants.
Essential workers had higher incidence than non-essential workers (IRR=1.65, 95% CI 1.102.26). Incidence rates were higher among participants from the South (IRR=1.67, 95% CI 1.08-2.59) compared to the Northeast U.S.

Table 3 shows the seroincidence and crude incidence rate ratios by epidemiologic risk 8 factors that were measured prior to serologic tests. Compared to those in single-family 9 dwellings with <4 household members, incidence was higher among those living in 10 multifamily dwellings with ≥4 household members (IRR=2.1, 95% CI 1.1-3.7) and those living 11 12 in congregate settings (IRR=2.5, 95% CI 1.2-4.8). A confirmed case in a household member was associated with a 15-fold higher incidence (IRR=16.3, 95% CI 9.6-27.8). Incidence was 13 higher among participants who dined indoors at restaurants or bars (IRR=1.93, 95% CI 1.39-14 2.70); visited a place of worship (IRR=1.92, 95% CI 1.26-2.84); gathered in groups ≥10 15 outdoors only (IRR=1.59, 95% CL1.07-2.34) as well as both indoors and outdoors (IRR=2.40 16 95% CI 1.41-3.09); visited indoors with non-household members sometimes wearing a mask 17 (IRR=1.79; 95% CI 1.11-2.96) or never wearing a mask (IRR=2.42; 95% CI 1.41-4.21); 18 worked indoors at a place of employment while never wearing a mask (IRR=2.45, 95% CI 19 0.96-5.34); wore masks only sometimes while attending a salon/gym (IRR=3.16, 95% CI 20 1.86-5.18); and reported travelling by air during August-November 2020 (IRR=1.50, 95% CI 21 1.04-2.14). 22

23 **Poisson models of SARS-CoV-2 seroconversion, May 2020-January 2021**

In crude analyses, participants who reported that they *always or sometimes* engaged in
social distancing with people they know (versus never) had a statistically significantly lower
seroincidence (IRR_{always vs never}=0.54, 95% CI 0.34-0.90; IRR_{sometimes vs never}=0.53, 95% CI 0.340.85). In multivariable analyses adjusted for sociodemographics and comorbidities (Model 1,
aIRR_{always vs never}=0.60, 95% CI 0.37-0.99), and additionally for community-level transmission

(Model 2, aIRR_{always vs never}=0.64, 95% CI 0.39-1.06), participants who reported always social 1 distancing with those they know (versus never) had lower seroincidence, although the 95% 2 3 confidence intervals were wider. Participants who reported social distancing always or 4 sometimes (vs. never) with people they did not know also had lower seroincidence rates in 5 both crude and adjusted models, but results were marginally significant. The composite risk score for SARS-CoV-2 incidence was associated with seroconversion in dose-response 6 7 fashion (IRR_{medium vs low} = 1.68, 95% CI 1.03-2.81; IRR_{high vs low} = 3.49, 95% CI 2.26-5.58) (Table 8 4). The sensitivity analysis that also included individuals who were seronegative at time 2, but 9 who reported a positive PCR or rapid test (n=187) did not materially alter the findings 10 11 (Supplemental Table S3). 12 Clinical and public health outcomes among persons with SARS-CoV-2 seroconversion during May 2020-January 2021 13 Among the 161 individuals who seroconverted during May 2020-January 2021, 97 (60.3%) 14 reported ever testing for SARS-CoV-2 outside the study, but only half (26.7% of total) 15 reported ever having a positive SARS-CoV-2 test. (Table 5). About 28% reported no 16 17 symptoms of COVID-like illness. Only 29.2% reported having ever isolated themselves from people outside their household because of their infection, and, among those who did not live 18 alone, even fewer (17.4% overall) reported having ever isolated themselves from others 19 20 within their household. Less than one-fifth (19.3%) of all seroconverters were asked about contacts following diagnosis and only 11.8% had been informed by a contact tracer about 21 contact with a confirmed SARS-CoV-2 case. Only 5.0% of all seroconverters were told by a 22 contact tracer to isolate because they had COVID-19. 23 DISCUSSION 24 25 We report findings from a large community-based prospective epidemiologic study of SARS-

CoV-2 incidence and risk factors in the U.S. during May 2020-January 2021. Using serologic
 tests, we longitudinally characterized the incidence of SARS-CoV-2 infection in relation to a
 range of risk factors. We found that social distancing and a low epidemiologic risk score

composed of modifiable risk factors was protective against infection, even after controlling
for other measured confounders. Finally, public health strategies such as quarantining,
testing, isolation, and contact tracing had low coverage and adoption among seroconverters,
limiting their effectiveness at reducing community transmission. Taken together, our study
findings document some of the principal reasons why the U.S. has continued to experience
sustained community transmission, hospitalizations, and deaths from COVID-19 in many
areas.

Social distancing was protective against seroincidence even after controlling for other risk 8 factors. This suggests a need for more effective and consistent messaging around social 9 10 distancing. We observed substantially increased risk for several other key epidemiologic risk factors reflected in a composite risk score. Among participants in the top tertile of the risk 11 12 score seroincidence risk was 3-fold higher, accounting for 55% of the observed seroconversions. Reducing multiple risk factors (e.g., through policies on masking, mass 13 gatherings, indoor dining/bars, social distancing, air travel) would likely substantially reduce 14 community transmission even in the vaccine era. 15

Our findings suggest that elevated risk among essential workers, observed early in the U.S. pandemic, persisted into the second phase of the pandemic. Essential workers risk exposure to SARS-CoV-2 not only in their workplaces, but also in their communities and as part of their work commute if they use public transportation. Household members of essential workers share their high infection risk [28]. Workplace safety measures, such mask/vaccine mandates, have the added benefit of protecting household members and other close contacts of essential workers.

We identified gaps in the reach of public health interventions aimed at reducing SARS-CoV-24 2 spread. Most who seroconverted did not report a prior positive PCR test, and a substantial 25 proportion were asymptomatic. Moreover, few people who seroconverted in our study 26 reported being reached by contact tracers. These results highlight the barriers to successful 27 implementation of isolation, contact tracing, and quarantine. Now that rapid home tests are

1 easily available, frequent proactive testing at home can be a more effective way to capture 2 asymptomatic and pre-symptomatic cases early and prevent onward transmission. 3 Our study highlights that the drivers of racial/ethnic disparities in SARS-CoV-2 risk need to 4 be targeted by governments, health departments and researchers [29]. Structural factors, 5 such as household crowding, the need to work in-person to avoid income loss, and 6 inequitable access to SARS-CoV-2 testing [30], create and perpetuate a disparate burden of 7 SARS-CoV-2 exposure and incidence [31]. To date, no targeted strategies or policies have 8 been deployed that aim to protect those who cannot afford missing work, including essential workers. Public health leaders and policy makers should proactively design pandemic 9 10 response strategies that counteract the prevailing structural forces, including structural 11 racism, that create, maintain, or exacerbate inequities in safety and health during a public 12 health crisis [32-35]. Our study has several limitations. The observed cumulative incidence may be lower than the 13

true cumulative incidence in our cohort because of waning of SARS-CoV-2 antibodies [36]. 14 Recent studies suggest waning of antibodies to both nucleocapsid and spike proteins [5]. 15 Combined with the timing of specimen collection relative to infection for many participants in 16 our cohort (median 190 days) [12], this could mean that we underestimated the true 17 cumulative incidence. The frequency of observed seroconversion in our cohort was 5% over 18 6 months (9.8 per 100 person-years), a level that is associated with slightly reduced positive 19 predictive value in single use assays, and may have resulted in misclassification of some 20 individuals as having seroconverted when they did not. This would bias estimated risk factor 21 22 associations toward the null. Next, estimated associations between SARS-CoV-2 risk factors and incidence are subject to confounding. The crude associations we presented may vary by 23 24 setting, with interpretation for some associations further hampered by small sample sizes. 25 The LASSO model is a predictive model, and the selected risk factors used in the composite risk score may not be causally associated with seroconversion. Some risk behaviors may 26 have been underreported, due to social desirability, which would bias observed associations 27 toward the null. Finally, our study period for the current analysis pre-dated the vaccine era 28

1 and the emergence of the highly transmissible and possibly more virulent Delta and Omicron

2 variants. We could not, therefore, examine risk factors for infection among vaccinated

3 persons. Lower vaccine effectiveness has been observed against the Delta and Omicron

4 variants for infection [37-41], and high viral loads have been documented among fully

5 vaccinated persons [3,42]. Thus, our findings related to transmission risk factors also likely

6 apply to vaccinated persons, as they remain at risk for breakthrough infection and onward

7 transmission of SARS-CoV-2 when engaging in some of the same risk factors.

8 Conclusion

9 Modifiable risk factors and poor reach of public health strategies continue to drive

10 transmission of SARS-CoV-2 across the U.S. While continuing to increase vaccine

11 coverage, it remains critical for public health agencies to simultaneously reduce risk factors

12 and address structural factors that contribute to high incidence and persistent inequities.

13 Future research will include monitoring SARS-CoV-2 outcomes in the vaccine, and Delta

14 and Omicron variant eras.

15

16 NOTES

17 ACKNOWLEDGEMENTS

The authors wish to thank the participants of the CHASING COVID Cohort Study. We are grateful to you for your contributions to the advancement of science around the SARS-CoV-2 pandemic. We thank Prof. Patrick Sullivan and MTL for local validation work on the serologic assays for use with DBS that greatly benefited our study. We are also grateful to MTL Labs for processing specimen collection kits and serologic testing of our cohort's specimens.

23 AUTHOR CONTRIBUTIONS

24 DN, MSR, SGK, MMR, conceptualized the study. MSR, MC and DN performed statistical

analyses. DN and MSR wrote the first draft of the paper. DN, MSR, MC, SGK, and AP

contributed to interpreting the data, DN, RZ, MSR, MC, SGK, WY, AB, CM, SK, AM, MMR,

27 DAW, AP, LW, and CG contributed to the writing and revising of the manuscript. SGK, WY,

- 1 AB, CM, SK, and DN contributed to data collection, cleaning and management. DN, SGK,
- 2 MMR and CG contributed to obtaining funding for the research.

3 FUNDING

- 4 This work was supported by The National Institute of Allergy and Infectious Diseases, award
- 5 number 3UH3AI133675-04S1 (MPIs: D Nash and C Grov); the CUNY Institute for
- 6 Implementation Science in Population Health (cunyisph.org); the COVID-19 Grant Program
- 7 of the CUNY Graduate School of Public Health and Health Policy; and the National Institute
- 8 of Child Health and Human Development grant P2C HD050924 (Carolina Population
- 9 Center).

10 **COMPETING INTEREST STATEMENT:** None to declare.

11

REFERENCES

2	1.	Buitrago-Garcia D, Egli-Gany D, Counotte MJ, et al. Occurrence and transmission
3		potential of asymptomatic and presymptomatic SARS-CoV-2 infections: A living
4		systematic review and meta-analysis. PLoS Med 2020; 17:e1003346.
5	2.	Kalish H, Klumpp-Thomas C, Hunsberger S, et al. Mapping a Pandemic: SARS-
6		CoV-2 Seropositivity in the United States. medRxiv 2021; Available at:
7		http://dx.doi.org/10.1101/2021.01.27.21250570.
8	3.	Brown CM, Vostok J, Johnson H, et al. Outbreak of SARS-CoV-2 Infections,
9		Including COVID-19 Vaccine Breakthrough Infections, Associated with Large Public
10		Gatherings — Barnstable County, Massachusetts, July 2021. MMWR. Morbidity
11		and Mortality Weekly Report. 2021; 70:1059–1062. Available at:
12		http://dx.doi.org/10.15585/mmwr.mm7031e2.
13	4.	Angulo FJ, Finelli L, Swerdlow DL. Estimation of US SARS-CoV-2 Infections,
14		Symptomatic Infections, Hospitalizations, and Deaths Using Seroprevalence
15		Surveys. JAMA Netw Open 2021 ; 4:e2033706.
16	5.	Bajema KL, Wiegand RE, Cuffe K, et al. Estimated SARS-CoV-2 Seroprevalence in
17		the US as of September 2020. JAMA Intern Med 2020; Available at:
18		http://dx.doi.org/10.1001/jamainternmed.2020.7976.
19	6.	CDC. COVID Data Tracker. 2020. Available at: https://covid.cdc.gov/covid-data-
20	(tracker/. Accessed 31 January 2021.
21	7.	Hutchins HJ, Wolff B, Leeb R, et al. COVID-19 Mitigation Behaviors by Age Group -
22		United States, April-June 2020. MMWR Morb Mortal Wkly Rep 2020; 69:1584–
23		1590.
24	8.	Oran DP, Topol EJ. The Proportion of SARS-CoV-2 Infections That Are
25		Asymptomatic : A Systematic Review. Ann Intern Med 2021; Available at:
26		http://dx.doi.org/10.7326/M20-6976.

1	1 9. Pouwels KB, House T, Pritchard E, et al. C	ommunity prevalence of SARS-CoV-2 in
2	2 England from April to November, 2020: res	ults from the ONS Coronavirus Infection
3	3 Survey. Lancet Public Health 2021 ; 6:e30–	e38.
4	4 10. Wei J, Stoesser N, Matthews PC, et al. Anti	body responses to SARS-CoV-2
5	5 vaccines in 45,965 adults from the general	population of the United Kingdom. Nat
6	6 Microbiol 2021 ; Available at: http://dx.doi.or	rg/10.1038/s41564-021-00947-3.
7	7 11. COVID-19 INFECTION SURVEY. Available	at: https://www.ndm.ox.ac.uk/covid-
8	8 19/covid-19-infection-survey. Accessed 1 A	August 2021.
9	9 12. Robertson M, Kulkarni S, Berry A, et al. A na	ational prospective cohort study of
10	10 SARS/COV2 pandemic outcomes in the U.	S.: The CHASING COVID Cohort.
11	11 Infectious Diseases (except HIV/AIDS). 202	20; Available at:
12	12 https://www.medrxiv.org/content/10.1101/2	020.04.28.20080630v1.
13	13 13. Nash D, Stief M, MacCrate C, et al. A Web-	Based Study of HIV Prevention in the
14	14 Era of Pre-Exposure Prophylaxis Among V	ulnerable HIV-Negative Gay and
15	15 Bisexual Men, Transmen, and Transwomer	n Who Have Sex With Men: Protocol for
16	16 an Observational Cohort Study. JMIR Rese	earch Protocols. 2019; 8:e13715.
17	17 Available at: http://dx.doi.org/10.2196/1371	5.
18	18 14. Grov C, Westmoreland D, Rendina HJ, Nas	h D. Seeing Is Believing? Unique
19	19 Capabilities of Internet-Only Studies as a T	ool for Implementation Research on HIV
20	20 Prevention for Men Who Have Sex With Me	en: A Review of Studies and
21	21 Methodological Considerations. J Acquir Im	nmune Defic Syndr 2019 ; 82 Suppl
22	22 3:S253–S260.	
23	23 15. Robertson MM, Kulkarni SG, Rane M, et al.	Cohort profile: a national, community-
24	24 based prospective cohort study of SARS-C	oV-2 pandemic outcomes in the USA-
25	25 the CHASING COVID Cohort study. BMJ C	0pen 2021 ; 11:e048778.
26	26 16. Center for Devices, Radiological Health. EU	A Authorized Serology Test
27	27 Performance. 2021. Available at: https://ww	vw.fda.gov/medical-devices/coronavirus-

1	disease-2019-covid-19-emergency-use-authorizations-medical-devices/eua-
2	authorized-serology-test-performance. Accessed 5 February 2021.
3	17. Tré-Hardy M, Wilmet A, Beukinga I, et al. Analytical and clinical validation of an
4	ELISA for specific SARS-CoV-2 IgG, IgA, and IgM antibodies. J Med Virol 2021;
5	93:803–811.
6	18. Trabaud M-A, Icard V, Milon M-P, Bal A, Lina B, Escuret V. Comparison of eight
7	commercial, high-throughput, automated or ELISA assays detecting SARS-CoV-2
8	IgG or total antibody. J Clin Virol 2020 ; 132:104613.
9	19. Plaga A, Wei R, Olson E, et al. Evaluation of the Clinical Performance of 7
10	Serological Assays for SARS-CoV-2 for Use in Clinical Laboratories. J Appl Lab
11	Med 2021 ; 6:998–1004.
12	20. Sempos CT, Tian L. Adjusting Coronavirus Prevalence Estimates for Laboratory
13	Test Kit Error. Am J Epidemiol 2021 ; 190:109–115.
14	21. Tibshirani R. Regression Shrinkage and Selection via the Lasso. J R Stat Soc
15	Series B Stat Methodol 1996; 58:267–288.
16	22. Wilson N, Kvalsvig A, Barnard LT, Baker MG. Case-fatality risk estimates for
17	COVID-19 calculated by using a lag time for fatality. Emerg Infect Dis 2020;
18	26:1339–1441.
19	23. Nash D, Qasmieh S, Robertson M, et al. Household factors and the risk of severe
20	COVID-like illness early in the US pandemic. medRxiv 2020; Available at:
21	http://dx.doi.org/10.1101/2020.12.03.20243683.
22	24. Lauer SA, Grantz KH, Bi Q, et al. The Incubation Period of Coronavirus Disease
23	2019 (COVID-19) From Publicly Reported Confirmed Cases: Estimation and
24	Application. Ann Intern Med 2020; 172:577–582.
25	25. Wang D, Hu B, Hu C, et al. Clinical Characteristics of 138 Hospitalized Patients With
26	2019 Novel Coronavirus–Infected Pneumonia in Wuhan, China. JAMA 2020;
27	323:1061–1069.

1	26. Russell TW, Hellewell J, Jarvis CI, et al. Estimating the infection and case fatality
2	ratio for coronavirus disease (COVID-19) using age-adjusted data from the
3	outbreak on the Diamond Princess cruise ship, February 2020. Eurosurveillance.
4	2020; 25. Available at: http://dx.doi.org/10.2807/1560-
5	7917.es.2020.25.12.2000256.
6	27. covid-19-data: An ongoing repository of data on coronavirus cases and deaths in the
7	U.S. Github, Available at: https://github.com/nytimes/covid-19-data. Accessed 18
8	August 2021.
9	28. Madewell ZJ, Yang Y, Longini IM Jr, Halloran ME, Dean NE. Household
10	Transmission of SARS-CoV-2: A Systematic Review and Meta-analysis. JAMA
11	Netw Open 2020 ; 3:e2031756.
12	29. Borrell LN, Elhawary JR, Fuentes-Afflick E, et al. Race and Genetic Ancestry in
13	Medicine - A Time for Reckoning with Racism. N Engl J Med 2021; 384:474–480.
14	30. Mody A, Pfeifauf K, Geng EH. Using Lorenz Curves to Measure Racial Inequities in
15	COVID-19 Testing. JAMA Netw Open 2021; 4:e2032696.
16	31. Ogedegbe G, Ravenell J, Adhikari S, et al. Assessment of Racial/Ethnic Disparities
17	in Hospitalization and Mortality in Patients With COVID-19 in New York City. JAMA
18	Netw Open 2020 ; 3:e2026881.
19	32. Nash D, Geng E. Goal-Aligned, Epidemic Intelligence for the Public Health
20	Response to the COVID-19 Pandemic. Am J Public Health 2020 ; 110:1154–1156.
21	33. Nash D. Designing and Disseminating Metrics to Support Jurisdictional Efforts to
22	End the Public Health Threat Posed by HIV Epidemics. Am J Public Health 2020;
23	110:53–57.
24	34. Bailey ZD, Feldman JM, Bassett MT. How Structural Racism Works - Racist Policies
25	as a Root Cause of U.S. Racial Health Inequities. N Engl J Med 2020; Available at:
26	http://dx.doi.org/10.1056/NEJMms2025396.

1	35. Bailey ZD, Krieger N, Agénor M, Graves J, Linos N, Bassett MT. Structural racism
2	and health inequities in the USA: evidence and interventions. Lancet 2017;
3	389:1453–1463.
4	36. Sullivan PS, Siegler AJ, Shioda K, et al. SARS-CoV-2 cumulative incidence, United
5	States, August-December 2020. Clin Infect Dis 2021; Available at:
6	http://dx.doi.org/10.1093/cid/ciab626.
7	37. Scobie HM, Johnson AG, Suthar AB, et al. Monitoring Incidence of COVID-19
8	Cases, Hospitalizations, and Deaths, by Vaccination Status - 13 U.S. Jurisdictions,
9	April 4-July 17, 2021. MMWR Morb Mortal Wkly Rep 2021; 70:1284–1290.
10	38. Rosenberg ES, Holtgrave DR, Dorabawila V, et al. New COVID-19 Cases and
11	Hospitalizations Among Adults, by Vaccination Status - New York, May 3-July 25,
12	2021. MMWR Morb Mortal Wkly Rep 2021 ; 70:1306–1311.
13	39. Cele S, Jackson L, Khan K, Khoury DS, Moyo-Gwete T, Tegally H, et al. SARS-
14	CoV-2 Omicron has extensive but incomplete escape of Pfizer BNT162b2 elicited
15	neutralization and requires ACE2 for infection. medRxiv. 2021 Dec
15 16	neutralization and requires ACE2 for infection. medRxiv. 2021 Dec 11;2021.12.08.21267417.
16	11;2021.12.08.21267417.
16 17	11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of
16 17 18	11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing
16 17 18 19	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805.
16 17 18 19 20	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805. Available from: <u>https://www.medrxiv.org/content/10.1101/2021.12.15.21267805v1</u>
16 17 18 19 20 21	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805. Available from: <u>https://www.medrxiv.org/content/10.1101/2021.12.15.21267805v1</u> 41. Lu L, Mok BW-Y, Chen L, Chan JM-C, Tsang OT-Y, Lam BH-S, et al. Neutralization
16 17 18 19 20 21 22	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805. Available from: <u>https://www.medrxiv.org/content/10.1101/2021.12.15.21267805v1</u> 41. Lu L, Mok BW-Y, Chen L, Chan JM-C, Tsang OT-Y, Lam BH-S, et al. Neutralization of SARS-CoV-2 Omicron variant by sera from BNT162b2 or Coronavac vaccine
16 17 18 19 20 21 22 23	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805. Available from: https://www.medrxiv.org/content/10.1101/2021.12.15.21267805v1 41. Lu L, Mok BW-Y, Chen L, Chan JM-C, Tsang OT-Y, Lam BH-S, et al. Neutralization of SARS-CoV-2 Omicron variant by sera from BNT162b2 or Coronavac vaccine recipients [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.13.21267668.
 16 17 18 19 20 21 22 23 24 	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805. Available from: https://www.medrxiv.org/content/10.1101/2021.12.15.21267805v1 41. Lu L, Mok BW-Y, Chen L, Chan JM-C, Tsang OT-Y, Lam BH-S, et al. Neutralization of SARS-CoV-2 Omicron variant by sera from BNT162b2 or Coronavac vaccine recipients [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.13.21267668. Available from: https://www.medrxiv.org/content/10.1101/2021.12.13.21267668.
 16 17 18 19 20 21 22 23 24 25 	 11;2021.12.08.21267417. 40. Doria-Rose N, Shen X, Schmidt SD, O'Dell S, McDanal C, Feng W, et al. Booster of mRNA-1273 Vaccine Reduces SARS-CoV-2 Omicron Escape from Neutralizing Antibodies [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.15.21267805. Available from: https://www.medrxiv.org/content/10.1101/2021.12.15.21267805v1 41. Lu L, Mok BW-Y, Chen L, Chan JM-C, Tsang OT-Y, Lam BH-S, et al. Neutralization of SARS-CoV-2 Omicron variant by sera from BNT162b2 or Coronavac vaccine recipients [Internet]. 2021 Dec [cited 2021 Dec 17] p. 2021.12.13.21267668. Available from: https://www.medrxiv.org/content/10.1101/2021.12.13.21267668v1 42. Chia PY, Xiang Ong SW, Chiew CJ, et al. Virological and serological kinetics of

1 **TABLES**

Table 1. Baseline characteristics of CHASING COVID Cohort Study participants who provided a dried blood spot sample for antibody testing

	Participants with one	Participants with one or more		
	serologic test	s		
	N	%		
Total	4,510	100.00		
Two serologic tests	3,605	79.93		
Age				
Median (IQR)	41 (31, 55)			
18-29	876	19.42		
30-39	1,253	27.78		
40-49	858	19.02		
50-59	658	14.59		
≥ 60	865	19.18		
Gender				
Male	2,018	44.75		
Female	2,360	52.33		
Non-Binary/Transgender	132	2.93		

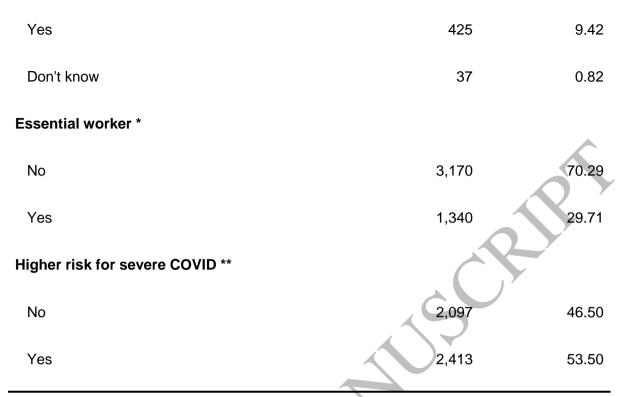
Race/Ethnicity

White non-Hispanic	2,966	65.76
Hispanic	679	15.06
Black non-Hispanic	378	8.38
Asian/Pacific Islander	304	6.74
Other	168	3.73
Missing	15	0.33
Education	S	
Less than high school	54	1.20
High school graduate	403	8.94
Some college	1,151	25.52
College graduate	2,902	64.35
Employment		
Employed	2,774	61.51
Out of work	544	12.06
Homemaker	242	5.37
Student	367	8.14
Retired	583	12.93
Household income		
<\$35,000	1 196	26.20

<\$35,000	1,186	26.30

\$35,000-49,999	496	11.00
\$50,000-69,999	643	14.26
\$70,000-99,999	741	16.43
≥\$100,000	1,329	29.47
Don't know	112	2.48
Missing	3	0.07
Setting		
Urban	1,911	42.37
Suburban	1,186	26.30
Rural	1,412	31.31
Missing	1	0.02
Geographic region		
Northeast	1,320	29.27
Midwest	805	17.85
South	1,269	28.14
West	1,111	24.63
US Territories	5	0.11
Healthcare worker		

No	4,048	89.76



* Combined from three follow-up interviews between August and November 2020

** >60 years old, or reported co-morbidity, or current smoker. Comorbidity was defined as having history of heart attack, depression, angina, immunosuppression, type 2 diabetes, high blood pressure, cancer, asthma, COPD, chronic kidney disease, and/or HIV/AIDS

	No. of seronegat ive participan ts in Period 1 *	No. of incident infection s in Period 2 **	Total person- years of follow up	Seroinciden ce per 100 person- years	Rate Ratio (95% CI)
Total	3,422	161	1,646.04	9.8 (8.3, 11.5)	Y
Age group				37	
			\sim	12.33 (8.9,	
18-29	617	37	300.00	16.7)	(ref)
				10.24 (7.6,	0.83 (0.53,
30-39	934	46	449.00	13.5)	1.28)
				10.65 (7.5,	0.86 (0.53,
40-49	654	33	310.00	14.7)	1.38)
				9.24 (6.1,	0.74 (0.43,
50-59	514	23	249.00	13.7)	1.25)
				6.49 (4.2,	0.52 (0.30,
≥ 60	703	22	339.00	9.8)	0.88)
Gender					
				11.76 (9.6,	
Male	1,516	88	748.00	14.3)	(ref)
				8.12 (6.4,	0.69 (0.50,
Female	1,810	69	850.00	10.2)	0.94)

Table 2: Crude associations of COVID-19 sociodemographic factors with seroincidence, May 2020-January 2021

Non-					
Binary/Transgend				8.21 (2.6,	0.69 (0.21,
er	96	4	48.71	20.6)	1.73)
Race/Ethnicity					
White non-				8.06 (6.5,	
Hispanic	2,307	92	1,141.00	9.8)	(ref)
				40.00/40.0	0.00 (4.44
Hisponia	500	37	219.00	16.89 (12.3,	2.09 (1.41,
Hispanic	500	57	219.00	22.6)	3.05)
Black non-				13.68 (8.2,	1.69 (0.96,
Hispanic	261	16	117.00	21.5)	2.82)
				6.56 (2.9,	0.81 (0.34,
Asian/PI	222	7	106.78	13.5)	1.66)
			K'		
				14.28 (7.1,	1.77 (0.84,
Other	132	9	63.02	25.9)	2.77)
Education					
Less than high				12.42 (2.1,	
school	37	2	16.10	39.4)	(ref)
High school				11.57 (6.7,	
graduate	278	14	121.00	18.9)	6.05)
				11.67 (8.7,	0.93 (0.27,
Some college	815	44	377.00	15.4)	5.76)
	5.0	••		,	5 0,
F				8.92 (7.3,	0.71 (0.21,
College graduate	2,292	101	1,132.00	10.8)	4.33)

Employment

				10.69 (8.8,	
Employed	2,102	109	1,020.00	12.8)	(ref)
				0.52 /5.0	0.00 (0.50
Out of work	402	18	188.90	9.53 (5.9, 14.9)	0.89 (0.52, 1.44)
	402	10	100.90	14.9)	1:44)
					0.86 (0.35,
Homemaker	179	7	78.70	8.89 (3.9, 18)	1.68)
Official	050	40	400.00	9.52 (5.2,	0.89 (0.46,
Student	253	12	126.00	16.4)	1.57)
				6.44 (3.7,	0.60 (0.33,
Retired	486	15	233.00	10.6)	1.01)
Household income					
				10.62 (7.8,	
<\$35,000	880	43	405.00	14.1)	(ref)
			×		. ,
		y		12.64 (8.3,	1.19 (0.70,
\$35,000-49,999	383	23	182.00	18.5)	1.96)
				7.47 (4.6,	0.70 (0.39,
\$50,000-69,999	505	18	241.00	11.7)	1.20)
		10	211100	,	0)
				11.89 (8.4,	1.12 (0.70,
\$70,000-99,999	592	34	286.00	16.3)	1.75)
				7.00/5.5	0 70 (0 40
≥\$100,000	993	38	495.00	7.68 (5.5, 10.5)	0.72 (0.46, 1.12)
20100,000	333	50	493.00	10.5)	1.12)
V				13.5 (5.1,	1.27 (0.44,
Don't know	77	5	37.00	29.5)	3.01)
Missing	2				

Setting

				8.7 (6.8,	
Urban	1,442	62	707.00	11.1)	(ref)
				9.58 (7.1,	1.11 (0.74,
Suburban	904	42	438.00	12.8)	1.65)
				11.37 (8.8,	1.29 (0.89,
Rural	1,076	57	501.00	14.6)	1.29)
Geographic region					
Northeast	977	34	477.00	7.13 (5, 9.9)	(ref)
				11.33 (8.1,	1.59 (0.98,
Midwest	629	34	300.00	15.6)	2.56)
				11.91 (9.1,	1.67 (1.08,
South	953	53	445.00	15.3)	2.59)
		*		9.44 (6.9,	1.32 (0.83,
West	859	39	413.00	12.8)	2.11)
U.S. Territories	4				
Healthcare worker					
No	3,074	140	1,481.00	9.45 (8, 11.1)	(ref)
				12.67 (7.9,	1.34 (0.80,
Yes	315	19	150.00	19.3)	2.12)
				14.3 (2.5,	1.51 (0.25,
Don't know	33	2	14.00	43.8)	5.07)

Essential worker

No	2,407	95	1,160.00	8.1 (6.7, 9.9)	(ref)
				13.6 (10.7,	1.65 (1.10,
Yes	1,015	66	486.00	17.0)	2.26)
High Risk group					5
				10.91 (8.8,	
No	1,574	83	761.00	13.4)	(ref)
				8.81 (7.1,	0.80 (0.59,
Yes	1,848	78	885.00	10.9)	1.10)
* May - September 2020					

** November 2020 - January 2021

*** Combined from three follow-up interviews between August and November 2020

**** >60 years old, or reported co-morbidity, or current smoker. Comorbidity was defined as having history of heart attack, depression, angina, immunosuppression, type 2 diabetes, high blood pressure, cancer, asthma, COPD, chronic kidney disease, and/or HIV/AIDS

1

Table 3: Crude associations of COVID-19 risk factors with seroincidence, May 2020-January 2021

	No. of seronegati ve participant s in Period 1 *	No. of incident infections in Period 2 **	Total person- years of follow up	Incidence rate per 100 person-years (95% CI)	Rate Ratio (95% CI)
Total	3,422	161	1,646.04	9.8 (8.3, 11.5)	
Household factors				2	
Multi-family with ≥4	. – .				
household members	171	14	78.00	17.9 (10.5, 28.6)	2.1 (1.1, 3.7)
Multi-family with <4					
household members	1,160	54	577.00	9.4 (7.2, 12.1)	1.1 (0.7, 1.6)
Single-family with ≥4		S'			
household members	625	27	289.00	9.3 (6.3, 13.4)	1.1 (0.7, 1.7)
Single-family with <4					
household members	1,374	57	660.00	8.6 (6.7, 11.1)	ref
Dorm, Group home, Other	92	9	42.04	21.4 (10.8, 37.2)	2.5 (1.2, 4.8)
Child in household	965	45	440.00	10.2 (7.6, 13.5)	1.1 (0.7, 1.5)
No child in household	2,457	116	1,206.00	9.6, 8.0, 11.5)	ref
Confirmed case in					
household member	28	15	123.6	12.1 (7.2, 19.6)	16.3 (9.6, 27.8)
No confirmed case in					
household member	3,394	146	19,630.80	0.7 (0.6, 0.9)	ref

Social distancing

Social distancing with people you do not know					
Always	2,616	113	1,258.00	8.9 (7.4, 10.7)	0.29 (0.14, 0.69)
Sometimes	660	34	319.00	10.6 (7.6, 14.7)	0.35 (0.16, 0.86)
Never	53	7	23.30	30.0 (13.8, 52.4)	ref
Not Applicable	53	4	24.50	16.3 (5.3, 37.5)	0.53 (0.13, 1.83)
				5	
Social distancing with people			\sim		
you know					
Always	1,137	51	543.00	9.4 (7.1, 12.2)	0.54 (0.33, 0.89)
Sometimes	1,787	79	868.00	9.1 (7.3, 11.3)	0.52 (0.33, 0.84)
Never	300	24	139.00	17.3 (11.5, 24.8)	ref
Not Applicable	158	4	74.50	5.4 (1.7, 13.9)	0.31 (0.09, 0.83)
	$\langle \mathbf{V} \rangle$				
Spent time in public places	Y				
Attended mass gathering(s)	350	18	174.89	10.3 (6.4, 16.5)	1.05 (0.63, 1.69)
Did not attend mass					
gathering(s)	3,072	143	1,471.15	9.7 (8.2, 11.5)	ref
Indoor dining/bar	1,755	108	843.86	12.8 (10.5, 15.5)	1.93 (1.39, 2.70)
No indoor dining/bar	1,667	53	802.18	6.6 (5.0, 8.7)	ref
Outdoor dining/bar	1,869	109	924.35	11.8 (9.8, 14.1)	1.1 (0.8, 1.4)

No outdoor dining/bar	1,553	78	721.70	10.8 (8.7, 13.3)	ref
Visited place of worship	359	29	168.87	17.2 (11.7, 25.1)	1.92 (1.26, 2.84)
Did not visit place of					
worship	3,063	132	1,477.17	8.9 (7.5, 10.6)	ref
Visited public park/public					
pool	2,334	100	1,147.00	8.7 (7.2, 10.5)	0.71 (0.52, 0.98)
Did not visit public					
park/pool	1,088	61	499.00	12.2 (9.5, 15.5)	ref
			A		
Gathered in groups ≥ 10		7			
No	1,961	70	939.00	7.4 (5.9, 9.4)	ref
Indoors only	230	11	109.00	10.1 (5.4, 7.7)	1.35 (0.68, 2.48)
Outdoors only	686	40	337.00	11.8 (8.7, 15.9)	1.59 (1.07, 2.34)
Indoors and outdoors	522	39	249.00	15.6 (11.4, 20.9)	2.40 (1.41, 3.09)
Do Not Know	23	1	11.70	8.5 (0.4, 41.0)	1.14 (0.05, 5.80)
	/				
Mask Use					
Mask while grocery shopping					
Did not go grocery					
shopping	173	9	84.60	10.6 (5.3, 19.7)	ref
Always	3,084	140	1,489.12	9.4 (7.9, 11.1)	0.88 (0.46, 1.84)
Sometimes	132	12	57.22	21.0 (11.6, 38.0)	1.97 (0.82, 4.86)
Never	33	0	15.11	0	0 (0, 2.21)

Mask while indoors visiting non-household members

Did no visit non-household					
members indoors	710	22	342.00	6.4 (4.1, 9.7) re	f
Always	1,118	43	546.26	7.9 (5.8, 10.7) 1.22 (0.73, 2.07)
Sometimes	1,131	63	546.44	11.5 (8.9, 14.9) 1.79 (1.11, 2.96)
				15.6 (10.9,	
Never	463	33	211.58	22.22) 2.42 (1.41, 4.21)
Mask while indoors at work			Δ)	
Did not attend indoor					
workplace	1,683	66	811.00	8.1 (6.4, 10.3) re	f
Always	1,372	71	660.60	10.7 (8.5, 13.6) 1.32 (0.94, 1.85)
Sometimes	299	18	144.08	12.5 (7.8, 20.1) 1.53 (0.88, 2.54)
Never	68	6	30.05	20.0 (8.6, 46.2) 2.45 (0.96, 5.34)
Mask while at salon/gym	,				
Did not attend salon/gym	1,607	59	771.00	7.6 (5.9, 9.8) re	f
Always	1,527	75	743.45	10.1 (8.0, 12.7) 1.31 (0.94, 1.86)
Sometimes	180	20	82.69	24.2 (15.2, 38.5) 3.16 (1.86, 5.18)
Never	108	7	48.94	14.3 (6.6, 30.8) 1.86 (0.78, 3.90)

Outdoor mask use

Mask use outdoors	1,562	67	753.27	8.9 (7.0, 11.4) 0.84 (0.67, 1.15)
-------------------	-------	----	--------	-----------------------------------

No mask use outdoors	1,860	94	892.78	10.5 (8.6, 13.0)	ref
Movement during the pandemic					•
Avoided public transit	2,692	124	1293	9.6 (8.1, 11.3)	ref
Used public transit	730	37	353	10.5 (7.6, 14.3)	1.09 (0.74, 1.56)
Recent Air travel (Aug-Nov)				R	
			Ċ		
Yes	582	39	287.91	13.5 (9.9, 18.2)	1.50 (1.04, 2.14)
No	2,840	122	1,358.00	8.9 (7.5, 10.6)	ref
Other potential risk factors		A			
Alcohol use ***	694	40	334.00	11.9 (8.8, 16.1)	1.50 (1.01, 2.22)
Substance use ****	914	45	443.40	10.1 (7.5, 13.7)	1.05 (0.73, 1.47)
Any comorbidities ^^	1,490	69	714.52	9.7 (7.6, 12.3)	0.97 (0.71, 1.33)
Changes in county-level	,				
community transmission					
High (0.327-3.30)	1,139	76	536.00 1	4.2 (11.4, 17.5)	ref
Medium (0.211-0.327)	1,140	37	557.00 6	.6 (4.8, 9.1)	0.46 (0.31, 0.69)
Low (0-0.211)	1,139	48	550.00 8	.7 (6.5, 11.4)	0.61 (0.42, 0.88)

Table 4: Crude and adjusted incidence rate ratios (IRRs) for seroincidence in the CHASING COVID Cohort Study, May 2020-January 2021

			Ad	justed	Ad	justed
	C	crude	(Model 1 *)		(Model 2 **)	
	IRR	95% CI	IRR	95% CI	IRR	95% Cl
Social distancing with people you						
do not know (ref:Never)						
Always	0.30	0.15, 0.72	0.37	0.18, 0.89	0.42	0.20, 1.00
Sometimes	0.35	0.16, 0.86	0.42	0.19, 1.05	0.47	0.22, 1.19
Not applicable (as per participant)	0.54	0.14, 1.80	0.70	0.18, 2.36	0.71	0.18, 2.38
Social distancing with people you						
do know (ref:Never)						
Always	0.54	0.34, 0.90	0.60	0.37, 0.99	0.64	0.39, 1.06
Sometimes	0.53	0.34, 0.85	0.57	0.36, 0.91	0.60	0.38, 0.96
Not applicable (as per participant)	0.31	0.09, 0.80	0.34	0.10, 0.90	0.37	0.10, 0.97
Composite measure of risk factors						
(ref:Low)						
Medium	1.59	0.98, 2.62	1.69	1.03, 2.81	1.68	1.03, 2.81
High	3.62	2.38, 5.71	3.53	2.29, 5.64	3.49	2.26, 5.58

* Model 1: Adjusted for age, gender, race/ethnicity, and comorbidities

** Model 2: Adjusted additionally for county-level changes in community level transmission

1

Table 5. Clinical and public health outcomes among persons with seroincident SARS-

CoV-2, May 2020-January 2021

	Ν	%
Total	161	100%
Symptoms and clinical outcomes		
PCR confirmed diagnosis	43	26.7
Asymptomatic *	45	28
Mild (symptomatic, but didn't seek care)	99	61.5
Ever had COVID like illness*	114	70.8
Nasal discharge, congestion or sneezing	100	62.1
Cough/Cough up phlegm	75	46.6
Cough up blood	0	0
Sore throat	65	40.4
Itchy eye or eye pain	53	32.9
Shortness of breath or chest pain	32	19.9
Stomachache, diarrhea, nausea or vomiting	67	41.6
Rash	12	7.5
Loss of smell	31	19.3
Headache	89	55.3

Fever, chills or repeated chills	44	27.3 2
Myalgia	58	3 36
Ever hospitalized	4	2.5
Public health outcomes and testing history		
Ever tested for COVID	97	60.3
Positive SARS-CoV-2 PCR test	43	26.7
Isolated from people outside household	47	29.2
Isolated from people within household **	28	17.4
Quarantined after contact with COVID	31	19.3
Asked about contacts after COVID diagnosis	31	19.3
Told about contacts with COVID case	19	11.8
Encouraged to get tested because of exposure to case	10	6.2
Told to stay home for a period of time	8	5

* Based on Council of State and Territorial Epidemiologists case definition

** Among those with others in the household