# Hybrid immunity from SARS-CoV-2 infection and vaccination in Canadian adults: cohort study

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- 15 REVISED May 28
- 16
- 17 Abstract: 314
- 18 Main text: 2269
- 19 4 figures and 1 table
- 20 Supplementary material: 5 figures and 1 table
- 21 References: 30

#### 22 Abstract

23 Background:

Few national-level studies have evaluated the impact of "hybrid" immunity (vaccination coupled with recovery from infection) from the Omicron variants of SARS-CoV-2.

26 27 Methods:

28 From May 2020 to December 2022, we conducted serial assessments (each of ~4000-9000 adults) examining

29 SARS-CoV-2 antibodies within a mostly representative Canadian cohort drawn from a national online polling

30 platform. Adults, most of whom were vaccinated, reported viral test–confirmed infections and mailed self-

31 collected dried blood spots to a central lab. Samples underwent highly sensitive and specific antibody assays to

32 spike and nucleocapsid protein antigens, the latter triggered only by infection. We estimated cumulative SARS-

33 CoV-2 incidence prior to the Omicron period and during the BA.1/1.1 and BA.2/5 waves. We assessed changes in

- 34 antibody levels and in age-specific active immunity levels.
- 35
- 36 Results:

37 Spike levels were higher in infected than in uninfected adults, regardless of vaccination doses. Among adults

38 vaccinated at least thrice and infected more than six months earlier, spike levels fell notably and continuously

39 for the nine months post-vaccination. By contrast, among adults infected within six months, spike levels

40 declined gradually. Declines were similar by sex, age group, and ethnicity. Recent vaccination attenuated

41 declines in spike levels from older infections. In a convenience sample, spike antibody and cellular responses

42 were correlated. Near the end of 2022, about 35% of adults above age 60 had their last vaccine dose more than

43 six months ago, and about 25% remained uninfected. The cumulative incidence of SARS-CoV-2 infection rose

44 from 13% (95% Cl 11-14%) before omicron to 78% (76-80%) by December 2022, equating to 25 million infected

45 adults cumulatively. However, the COVID-19 weekly death rate during the BA.2/5 waves was less than half of

46 that during the BA.1/1.1 wave, implying a protective role for hybrid immunity.

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48 Conclusions:

49 Strategies to maintain population-level hybrid immunity require up-to-date vaccination coverage, including

50 among those recovering from infection. Population-based, self-collected dried blood spots are a practicable

- 51 biological surveillance platform.
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53 Funding:

54 Funding was provided by the COVID-19 Immunity Task Force, Canadian Institutes of Health Research, Pfizer

55 Global Medical Grants, and St. Michael's Hospital Foundation. PJ and ACG are funded by the Canada Research

- 56 Chairs Program.
- 57

#### 58 Introduction

Infection with the Omicron BA.1/1.1 variant of the SARS-CoV-2 virus occurred worldwide late in 2021 and in early 2022. "Hybrid" immunity (vaccination coupled with recovery from infection) has emerged as a major determinant of the lower burden of COVID-19 morbidity and mortality in 2022 compared to 2020 or 2021, and as a key determinant of current population-based immunity.<sup>1,2</sup>

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Epidemiological studies have identified hybrid immunity as partially protective against infection or reinfection, and more strongly protective against hospitalization, severe disease, or death.<sup>1-3</sup> However, such studies rely on the follow-up of hospitalized patients or those with access to PCR-based testing, and not randomly selected populations. Thus, the contribution of infection and vaccination to hybrid immunity and the duration of immunity from either exposure remain remarkably poorly documented at the population level.<sup>4-7</sup>

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70 Development of strategies to move from pandemic to endemic management of COVID-19 will be greatly

71 enabled by evidence of population-level immunity, which ideally should be informed by changes over time in

biologic measures of immunologic protection (antibody levels, infection status, vaccination, and healthcare
 utilization). Humoral antibody levels, which correlate strongly with cellular immunity,<sup>8</sup> are the most practical
 method to monitor populations.

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Canada provides an opportunity to document hybrid immunity. Although reaching high levels of vaccination
 reasonably quickly (by September 2021), Canada experienced a large increase in infections from Omicron from
 December 2021, even among vaccinated people.<sup>9</sup> Vaccines used in Canada (mostly the mRNA and some
 adenovirus vaccines) trigger antibody responses to the SARS-CoV-2 spike protein and its receptor-binding
 domain (RBD), but not to the nucleocapsid protein (N).<sup>10</sup> This enables serological distinction of infection from
 vaccination.

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In this study, we estimate cumulative SARS-CoV-2 incidence among Canadian adults in 2020<sup>4</sup> and 2021 - prior to
 the Omicron period - and during two major Omicron waves (BA.1/1.1 and BA.2 and BA.5) in 2022.<sup>5</sup> We assess
 declines in active immunity and changes over time in age-specific active immunity levels based on prior infection
 and concurrent vaccination.

#### 88 Methods

89 From May 2020, the Action to Beat Coronavirus (Ab-C) study conducted six serial assessments of SARS-CoV-2 90 symptoms (via online surveys) and seropositivity (via antibody testing), with five surveys covering about 4000-91 9000 adults (fig 1). We recruited adults using the Angus Reid Forum, a nationally representative online polling 92 platform that approximately matches Canada's demographic profile.<sup>4</sup> We obtained informed consent from each 93 participant and excluded any unconsented panelist from the study. Details of the sampling, antibody testing strategy, and analyses have been published earlier.<sup>4,5,11,12</sup> The supplementary methods and Appendix 2-figure 1 94 95 report the recruitment, the dried blood spot (DBS) sample return rates, and the few exclusions from the six 96 phases of the study.

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98 The online survey assessed demographic characteristics, history of smoking, hypertension, obesity (self-reported 99 height and weight), diabetes, experience with SARS-CoV-2 infection symptoms, and COVID-19 testing (PCR or

rapid antigen). At the end of the survey, respondents indicated their willingness to provide a blood sample by

101 finger prick, and we sent consenters a DBS collection kit with instructions to self-collect. DBS samples were

- 102 returned to Unity Health laboratories in Toronto, with mail transit times ranging 3-6 days. Sinai Health in
- 103 Toronto conducted highly sensitive and specific chemiluminescence-based enzyme-linked immunosorbent
- assays targeting the spike protein, RBD, and N; validation of the assays is reported elsewhere.<sup>13,14</sup> Various quality
- 105 control steps focused on reducing false positives and false negatives, as well as adjusting the dilution to better 106 detect antibody signals after vaccination became widespread (Appendix 1 provides details of the lab methods
- and analyses). We conducted cluster analyses of N-positivity (defined below) to assign a probability of
- seropositivity to each sample using control samples and those with known past viral testing results (Appendix 2-
- 109 figure 2). In a subset of 39 adults in Toronto selected conveniently, we collected venous blood samples at home,
- 110 and tested these centrally for cellular immunity using the Euroimmun Interferon Gamma Release Assay<sup>15</sup> to
- 111 detect T-cell activity against the spike protein (supplementary methods).
- 112
- 113 Our primary outcomes were the relative levels of antibodies to the spike protein (hereafter "spike levels"),
- 114 which are increased both by vaccination and infection (defined as N-positivity or self-reported PCR/rapid test
- positivity), as a proxy for hybrid immunity levels. Our secondary outcome was the combination of vaccination
- 116 history and infection. We applied the age-specific cumulative incidence of SARS-CoV-2 to the Statistics Canada
- 117 national population totals<sup>16</sup> to derive estimates of the number of adults infected in each major phase and
- 118 compared cumulative incidence to confirmed COVID deaths by phase. Confirmed COVID deaths in Canada<sup>9</sup> are
- within 10% of analyses that apply excess all-cause mortality as an upper bound for COVID-19 mortality.<sup>17</sup>

#### 121 Results

We examined three time periods: (i) March 2020 to December 2021 when Canada faced waves of ancestral, Alpha, and Delta variants of SARS-CoV-2; (ii) January-March 2022 during the Omicron BA.1/1.1 wave; and (iii) April-December 2022 during the Omicron BA.2 and BA.5 waves. Figure 1 provides the timeline for Phases 1 to 6, in relation to national weekly averages of confirmed COVID-19 cases and weekly averages of vaccination from any dose.

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128 We surveyed 10 088 adults in Phase 6 of Ab-C, of whom 4025 provided DBS from 26 September to 21 November 129 2022, and of whom 3378 provided both surveys and DBS. Study participants were comparable to Canadian 130 adults in prevalence of obesity, smoking, diabetes, and vaccination, but fewer lower-education adults 131 participated (supplementary table S1). More females and vaccinated adults provided DBS in Phase 6. Lack of 132 vaccination and lower education were correlated (supplementary methods), so we adjusted cumulative 133 incidence for vaccination status. The characteristics of the cohort changed little between Phases 3, 4, and 6 134 (Appendix 3-table 1), so changes in antibody levels are unlikely to be confounded by differential recruitment in each phase.4,5 135

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137 Canada had four major viral waves before December 2021 and a major increase in vaccination coverage with 138 two doses peaking in early July 2021 (fig 1). A large Omicron BA.1/1.1 wave of January-March 2022 coincided 139 with a large increase in vaccination, mostly of third (booster) doses. The six Ab-C phases captured Canada's 140 major infection and vaccination peaks in a reasonably timely manner.

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Spike levels were higher in infected than in uninfected adults, regardless of vaccination doses (fig 2). Spike levels were higher among those who were infected and vaccinated, and lowest among the very few who remained

- 144 uninfected and unvaccinated, or had only one vaccine dose, or infection without vaccination. Uninfected adults
  - 4

with four vaccine doses were similar in spike level distribution to infected adults with only two or three vaccinedoses. Results using the RBD protein were similar (Appendix 2-figure 3).

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148 Among adults vaccinated at least thrice and infected more than six months prior to the last vaccine dose, spike 149 levels fell notably and continuously for the nine months post-vaccination (fig 3). By contrast, among adults 150 infected within six months, the decline in spike levels was more gradual. Declines were similar by sex, by age 151 group (15-59 years or 60+ years), and among various ethnicities (including visible minorities and Indigenous 152 populations). Vaccination within six months boosted spike levels from older infections that would have 153 otherwise fallen, yielding similar spike levels among adults infected more than six months ago or infected within 154 six months (supplementary figure S4). Stratifying by periods of 2 months or less, 3-5 months, and 6 or more 155 months yielded comparable results, albeit with smaller numbers in each stratum (data not shown).

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Among a convenience sample of 39 adults, all 32 vaccinated adults had positive spike T-cell responses. The T-cell
 titers and spike antibody levels correlated (Appendix 2-figure 5).

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160 Applying the Ab-C results, after standardizing for vaccination status, region, age, and sex to the national profile 161 of Canadian adult population, yielded estimates of cumulative incidence of SARS-CoV-2 infection rising from 162 about 13% before Omicron to 78% by December 2022. This equates to about 25 million infected adults 163 cumulatively. Canada had about 50 000 COVID deaths from March 2020 to December 2022, corresponding to about 6% higher mortality at all ages versus background death rates.<sup>17</sup> Over 90% of Canadian COVID deaths 164 occurred above age 60 years.<sup>9</sup> Despite the rising cumulative incidence, the COVID-19 weekly death rate per 165 166 million population during the Omicron BA.2/5 waves (7.7) was less than half of the weekly death rate during the 167 Omicron BA.1/1.1 wave (16.6). This suggests that hybrid immunity played a role in reducing severe disease and

168 deaths (Table 1), at least prior to the eventual waning of the immunity.<sup>18,19</sup>

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170 There were marked increases in infection among younger (18-59 years) and older (60+ years) mostly vaccinated 171 adults, rising from about 11% in each age group by August 2021 to about 86% and 75%, respectively, by

172 December 2022 (fig 4). However, fully 35% of adults above age 60, who are most at risk of hospitalization or

173 death, had their last vaccine dose more than six months ago, and about 25% remained uninfected.

174175 Discussion

We demonstrate the protective nature of hybrid immunity at a population level using robust biological markers of cumulative infection paired with viral testing. While steps to protect individuals and populations from SARS-CoV-2 infection must continue to be implemented, close to 80% of Canadian adults became infected, mostly from the Omicron variants, by December 2022. This high level of infection from the Omicron variants not only led to notable morbidity and mortality, but also contributed to population-level hybrid immunity.

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182 Despite a marked increase in cumulative infection, COVID-19 death rates during Omicron BA.2 and BA.5 were

183 markedly lower than during BA.1/1.1, likely reflecting a strong correlation between protection against severe

184 disease and hybrid immunity (despite lower protection against reinfection). Canadian healthcare systems were 185 overburdened with COVID-related hospitalizations several times during the pandemic. Since summer 2022,

- hospitalizations have eased significantly, most notably with fewer admissions to intensive care units following
- 187 the initial Omicron BA.1/1.1 wave.<sup>9</sup> Continued COVID-related practices (most of which were dropped on October
- 188 1, 2022), such as travel restrictions, masking mandates, and testing requirements, also may have played a role in

the lessened severity of COVID outcomes. Differences in pathogenicity of successive Omicron variants are likely
 too small<sup>20</sup> to explain the differences in COVID-19 death rates.

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192 We showed that absent recent infection, spike levels declined up to nine months, but reassuringly, declines 193 were comparable in older versus younger adults and by sex and ethnicity. Importantly, recent vaccination 194 attenuated the declines in spike levels from older infections. Obviously, reliance on infections is unwise to boost 195 immunity, especially for those most vulnerable to severe COVID-19. Collectively, our and other studies on hybrid 196 immunity<sup>1-5,7,21</sup> suggest that older adults may require access to booster doses at 6 to 12-month intervals and 197 prior to possible seasonal waves to achieve a robust level of protection against infection. Strategies to maintain 198 population-level hybrid immunity require high vaccination coverage, including among those who have recovered 199 from infection and the few remaining unvaccinated.

200

201 The Ab-C study is one of the few nationally representative serosurveys to measure hybrid immunity objectively,<sup>5-7</sup> and has the benefit of sampling the entire population. Large increases from Omicron wave are 202 evident in other Canadian studies (mostly done prior to the BA.5 waves).<sup>22</sup> A national US study among blood 203 204 donors reports lower levels of infection than do we,<sup>6</sup> but has not yet reported on the BA.4/5 waves. Moreover, blood donors or hospitalized patients may have notable biases.<sup>22</sup> Since the Omicron variant of SARS-CoV-2 205 206 appeared, self-testing using rapid antigen tests displaced PCR-testing in many countries, including Canada.<sup>23</sup> The 207 use of spike levels has limitations, although we found it correlated with cellular immunity. Earlier studies demonstrate that high levels of spike or RBD antibodies are predictive of neutralizing antibodies<sup>8</sup> and correlate 208 with lower viral loads that reduce severe disease in the infected and transmission to others.<sup>24</sup> 209

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211 Nonetheless, our study has some limitations. First, we had a larger proportion of highly educated adults than the 212 Canadian population. However, the selection biases did not change with subsequent waves, and we saw 213 widespread infection and vaccination in all education groups. We deliberately focused on distributions of 214 antibody levels which overlap in the comparison categories, but this has the benefit of showing the full range of spike antibody response in the various strata of the infected and vaccinated. We may be underestimating spike 215 antibody levels due to assay saturation.<sup>13</sup> N-positivity may have underestimated actual infection because mild 216 217 cases among vaccinated adults did not mount an antibody response or because people did not seroconvert 218 during the sampling period. Conversely, some adults may have reverted to N-negative status. Finally, defining 219 infection based on cumulative seropositivity and time-specific viral test positivity is crude and made more 220 complicated by periodic viral or vaccination waves. Thus, we are limited in quantifying the hybrid immunity 221 arising from various sequences of variant infections and vaccinations. For example, the apparent plateauing of 222 spike level declines at nine months in figure 3 may reflect cohorts facing at least two distinct vaccination or viral 223 waves. Future phases of our study may assess long-term immunity across different populations, as well as 224 development of long COVID.

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226 Canadian COVID-19 death rates are lower compared to the United States and other similar countries,<sup>25</sup> and we 227 speculate this may be from the sequence of low levels of infection pre-Omicron paired with high vaccination 228 coverage of two doses, followed by a large Omicron wave. Comparative analyses across countries using 229 objective measures of hybrid immunity are required. In Canada and other countries, home-based self-drawn 230 dried blood spots are a widely practicable and relatively inexpensive monitoring strategy for SARS-CoV-2 231 population immunity. Despite their limitations, serial serosurveys at the population level are reasonably 232 efficient, low-cost ways to monitor hybrid immunity and to study newer variants of SARS-CoV-2, and possibly 233 even other infectious agents. Future directions could include routine monitoring of various respiratory 234 pathogens, and work to develop practicable multi-plex assays for such infections.

#### 235 Acknowledgements

- 236 We thank the thousands of Canadians who participated in the Ab-C study. Euroimmun Medical Diagnostics (Sean
- 237 McFadden) supported the T-cell testing platform at St. Joseph's Health Centre/Unity Health. We thank the
- thousands of Canadians who participated in the Action to Beat Coronavirus study. A full listing for the Ab-C
- 239 Study Collaborators is available at <u>www.abcstudy.ca</u>.

#### 241 Competing interests

- ASS has received consulting fees from Apeiron Biologics, Cellenkos, Diffusion Pharmaceuticals, and
- GlaxoSmithKline outside the submitted work. IIB has served as a consultant for BlueDot and the National Hockey
   League Players' Association outside the submitted work.

#### 246 Funding

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Funding was provided by the COVID-19 Immunity Task Force, Canadian Institutes of Health Research, Pfizer Global Medical Grants, and St. Michael's Hospital Foundation. PJ and ACG are funded by the Canada Research Chairs Program. The funders had no role in study design, data collection and interpretation, or the decision to submit the work for publication.

#### 252 Ethics

The Ab-C study was approved by the Unity Health Toronto Research Ethics Board (REB # 20-107 and 21-213). All participants provided informed consent to be included in the study.

#### 256 Data availability

- Ab-C data will be made available publicly through the COVID-19 Immunity Task Force (CITF) Databank. To access the data, please create an account on the CITF Databank portal and submit an application to use the data. Your application will be reviewed by the CITF Databank team. The data access procedure is described in detail at <u>https://www.covid19immunitytaskforce.ca/wp-content/uploads/2022/11/data-access-diagram-en.pdf</u>. This process is free of charge.
- 262
- Analytical code will be available on request in accordance with the Ab-C study's data governance plan. Please email the corresponding author, Dr. Jha at <u>prabhat.jha@utoronto.ca</u> to request the code. The CITF data team harmonizes data from multiple studies funded by CITF, including the Ab-C study. As a result, variable names and labels may change after the harmonization. To minimize confusion when using the code, it's best to have some
- 267 contact with us when using the harmonized data.
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#### 269 References

- 270 1 Bobrovitz N, Ware H, Ma X, et al. Protective effectiveness of previous SARS-CoV-2 infection and hybrid
- immunity against the omicron variant and severe disease: a systematic review and meta-regression. *Lancet Infect Dis* 2023;S1473-3099(22)00801-5.
- 273 2 COVID-19 Forecasting Team. Past SARS-CoV-2 infection protection against re-infection: a systematic review 274 and meta-analysis. *Lancet* 2022;401:833-42.
- Altarawneh HN, Chemaitelly H, Hasan MR, et al. Protection against the Omicron variant from previous SARS-CoV-2 infection. *N Engl J Med* 2022;386:1288-90.
- Tang X, Sharma A, Pasic M, et al. Assessment of SARS-CoV-2 seropositivity during the first and second viral
  waves in 2020 and 2021 among Canadian adults. *JAMA Netw Open* 2022;5:e2146798.
- Brown PE, Fu SH, Bansal A, et al. Omicron BA.1/1.1 SARS-CoV-2 infection among vaccinated Canadian adults.
   *N Engl J Med* 2022;386:2337-9.
- Centers for Disease Control and Prevention. COVID Data Tracker. Atlanta, GA: US Department of Health and
   Human Services, CDC, 2023. <u>https://covid.cdc.gov/covid-data-tracker</u> (accessed 28 February 2023).
- Goldberg Y, Mandel M, Bar-On YM, et al. Protection and waning of natural and hybrid immunity to SARS CoV-2. *N Engl J Med* 2022;386:2201-12.
- Feng C, Shi J, Fan Q, et al. Protective humoral and cellular immune responses to SARS-CoV-2 persist up to 1
   year after recovery. *Nat Commun* 2021;12:4984.
- Public Health Agency of Canada. Coronavirus disease (COVID-19). Ottawa, ON: Public Health Agency of
   Canada, 2023. <u>https://health-infobase.canada.ca/covid-19/</u> (accessed 28 February 2023).
- 10 Duarte N, Yanes-Lane M, Arora RK, et al. Adapting serosurveys for the SARS-CoV-2 vaccine era. *Open Forum* 290 *Infect Dis* 2022;9:ofab632.
- 11 Wu DC, Jha P, Lam T, et al. Predictors of self-reported symptoms and testing for COVID-19 in Canada using a
   nationally representative survey. *PLoS One* 2020;15:e0240778.
- 293 12 Tang X, Gelband H, Nagelkerke N, et al. COVID-19 vaccination intention during early vaccine rollout in
- Canada: a nationwide online survey. *Lancet Reg Health Am* 2021;2:100055.
- Colwill K, Galipeau Y, Stuible M, et al. A scalable serology solution for profiling humoral immune responses
   to SARS-CoV-2 infection and vaccination. *Clin Transl Immunology* 2022;11:e1380.
- 14 Isho B, Abe KT, Zuo M, et al. Persistence of serum and saliva antibody responses to SARS-CoV-2 spike
   antigens in COVID-19 patients. *Sci Immunol* 2020;5:eabe5511.
- 299 15 Fernandez-Gonzalez M, Agullo V, Padilla S, et al. Clinical performance of a standardized Severe Acute
- Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) interferon-gamma release assay for simple detection of T-cell responses after infection or vaccination. *Clin Infect Dis* 2022;75:e338-46.
- 302 16 Statistics Canada. Table 17-10-0005-01 population estimates on July 1st, by age and sex. 2023.
- 303 <u>https://doi.org/10.25318/1710000501-eng</u> (accessed 15 February 2023).
- 304 17 World Health Organization. Global excess deaths associated with COVID-19 (modelled estimates). Geneva:

World Health Organization, 2022. <u>https://www.who.int/data/sets/global-excess-deaths-associated-with-covid-</u>
 19-modelled-estimates (accessed 28 February 2023).

- 307 18 Chemaitelly H, Nagelkerke N, Ayoub HH, et al. Duration of immune protection of SARS-CoV-2 natural 308 infection against reinfection. *J Travel Med* 2022;29:taac109.
- 309 19 Chemaitelly H, Tang P, Hasan MR, et al. Waning of BNT162b2 vaccine protection against SARS-CoV-2
- 310 infection in Qatar. *N Engl J Med* 2021;385:e83.
- 20 Strasser ZH, Greifer N, Hadavand A, Murphy SN, Estiri H. Estimates of SARS-CoV-2 Omicron BA.2 subvariant
- 312 severity in New England. *JAMA Netw Open* 2022;5:e2238354.
- 21 Tan CY, Chiew CJ, Lee VJ, Ong B, Lye DC, Tan KB. Comparative effectiveness of 3 or 4 doses of mRNA and
- 314 inactivated whole-virus vaccines against COVID-19 infection, hospitalization and severe outcomes among elderly
- in Singapore. *Lancet Reg Health West Pac* 2022;29:100654.

- 22 Murphy TJ, Swail H, Jain J, et al. The evolution of SARS-CoV-2 seroprevalence in Canada: a time-series study,
- 317 2020–2023. Can Med Assoc J 2023;195:E1030-E1037.
- 318 23 Angus Reid Forum. Incidence of Omicron: One-in-five Canadians report COVID-19 infection in their
- household since Dec. 1. Angus Reid Forum, 2022. <u>https://angusreid.org/omicron-incidence-restrictions/</u>
   (accessed 28 February 2023).
- 321 24 Tan ST, Kwan AT, Rodriguez-Barraquer I, et al. Infectiousness of SARS-CoV-2 breakthrough infections and
- reinfections during the Omicron wave. *Nat Med* 2023;29(2):358-65.
- Razak F, Shin S, Naylor CD, Slutsky AS. Canada's response to the initial 2 years of the COVID-19 pandemic: a comparison with peer countries. *Can Med Assoc J* 2022;194:E870-7.
- 26 Little N. COVID-19 Tracker Canada. <u>https://covid19tracker.ca/</u> (accessed 3 February 2023).
- 326 27 Fox J, Weisberg S. An R companion to applied regression. Thousand Oaks, CA: Sage Publications; 2018.
- 327 28 Angus Reid Institute. How we poll. http://angusreid.org/how-we-poll-ari/ (accessed 17 December 2020).
- 328 29 Action to Beat Coronavirus Study. Participant Information Sheet.
- 329 https://abcstudy.ca/docs/abcstudy\_information.pdf (accessed 25 November 2021).
- 30 Jackson C. delta method: The delta method. In: msm: Multi-State Markov and Hidden Markov Models in
- 331 Continuous Time. https://rdrr.io/cran/msm/man/deltamethod.html (accessed 3 February 2023).

## 333 Table 1. Cumulative incidence, numbers of infected adults, cumulative deaths, and period

## 334 COVID-19 mortality rate in Canada during various SARS-CoV-2 viral waves

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Time period	Cumulative incidence* % (95% CI)	No of adult (age 18 or older) infections in millions	Cumulative no of deaths†	Covid-19 mortality rate per million per week during the relevant period		
Pre-Omicron 2020-2021	12.7 (11.2-14.1)	3.9 (3.5-4.4)	30 1 4 9	8.6		
Omicron BA.1/1.1 JanMar. 2022	35.7 (34.0-37.4)	11.3 (10.7-11.8)	37 750	16.6		
Omicron BA.2/5 AprDec. 2022	77.7 (75.7-79.6)	24.6 (23.9-25.2)	49 674	7.7		

336 Notes:

337 \*Post-stratified for geographic region, age, sex, and vaccination status to derive the mean estimate (supplementary methods).

338 †We used data by end of December 2021, March 2022, and December 2022 from Public Health Agency of Canada's COVID-19 epidemiology

339 update (<u>https://health-infobase.canada.ca/covid-19/</u>) for total number of deaths.<sup>9</sup> Applying the proportion of long-term care deaths from Long-term

340 Care COVID-19 Tracker (https://ltc-covid19-tracker.ca) to the last period, 19789 of total cumulative deaths occurred in long-term care. Of all long-

341 term care deaths, about 80% occurred during the pre-Omicron period, mostly during the first viral wave of March-June 2020 (fig 1). Over 90% of

all COVID deaths occurred at ages 60 or older.

344 Figure 1. Seven-day rolling averages of PCR-confirmed COVID-19 cases in

345 Canada (black solid and dotted line), and SARS-CoV-2 vaccinations (any dose;

red line) in relation to the data collection phases of the Ab-C study. Testing and

- vaccination data were derived from COVID-19 Tracker Canada as of 3 February 2023
- 348 (https://COVID19Tracker.ca).<sup>26</sup> Data on major variants were obtained from Public
- Health Agency of Canada's Health Infobase COVID-19 epidemiology update
- 350 (https://health-infobase.canada.ca/covid-19/testing-variants.html).<sup>9</sup> Dotted lines for
- PCR-based testing after 1 January 2022 reflect the major uncertainty in PCR-based
   testing. Widespread PCR testing guidelines became stricter and were significantly
- scaled back in community settings and thus became far less reliable to monitor trends.
- 353 E 354

## Figure 2. Levels of antibodies to the spike protein stratified by infection and number of vaccination doses.

- 357 Circles represent individuals with their last vaccination (or unvaccinated) >10 days prior
- to dried blood spot sample collection (n=3378 with complete information available as of
- the time of analyses after excluding 14 low quality samples). We further excluded 16
- 360 participants whose samples were seronegative and viral test was positive, but who did
- 361 not provide viral test dates or reported test dates less than eight days from the receipt of
- 362 DBS. The solid-coloured line represents the median and box plots show the interquartile
- range. The results above a relative level of 1.2 are outside the linear range of the assay.
   Results using the receptor-binding domain antigen were similar to the spike protein
- 365 (supplementary figure S3).
- 366

### 367 Figure 3. Age, sex, and ethnicity-specific trends to nine months in levels of

- antibodies to the spike protein among adults vaccinated with 3-4 doses, stratified
   by infection more than six months ago or less than six months ago.
- 370 See footnote to figure 2 for testing details. We created smoothed curves and 95%
- 371 confidence intervals using locally weighted scatterplot smoothing with span parameter
   372 of 0.8.<sup>27</sup>
- 373

### **Figure 4. Cumulative incidence in each stratum of infection and vaccination in the**

- pre-Omicron wave, during the Omicron BA.1/1.1 wave, and during the BA.2 and
  BA.5 waves by age group.
- \*Including uninfected and infected cases. The first column in each age group represents
  the antibody and viral test positivity for the entire period prior to Omicron, whereas the
  second column represents the values during the omicron BA.1/1.1 wave and the third
- during the BA.2/5 waves. By the last time period studied, the numbers of participants
- aged 15-59 who were N-positive, viral test–positive, and positive to both were 675
- 382 (41%), 37 (2%), and 699 (43%). The comparable numbers for participants aged 60 or
- 383 more were 763 (44%), 35 (2%), and 500 (29%).

- **Supplementary material** Brown PE, Fu SH, Newcombe L, et al. Hybrid immunity from SARS-CoV-2 infection and vaccination 385
- in Canadian adults: cohort study. 386

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#### 404 Methods

#### 405 Subject recruitment

406 The Action to Beat Coronavirus (Ab-C) study received ethical approval from Unity Health Toronto (REB 20-107). In Phase 407 1, from May through September 2020, we invited 44 270 members (out of about 78 000 total members) of the Angus Reid 408 Forum,<sup>28</sup> an established nationwide polling panel of Canadian adults aged 18 and older, to complete an online survey about 409 SARS-CoV-2 symptoms and testing histories. The sampled population was stratified by age groups (18-34, 35-54, 55+); sex 410 (male, female); education (high school education or lower, some college or college or technical degree, some university, or 411 university degree); and region, by census metropolitan area to match the national demographic profile, with oversampling of 412 adults 60 years or older. In August 2021, we invited about 3100 additional Forum panel members from 17 regions with high 413 burden of infection (of 93 total regions nationwide), based on a regression analysis of SARS-CoV-2 case counts.<sup>4</sup> From 414 December 2020 through January 2021, we invited all 19 994 Phase 1 participants to join Phase 2, retaining the same sampling 415 frame. Phase 3 and 4 recruitment used similar approaches. In Phase 4, we conducted additional outreach to 2587 additional 416 members from marginalized groups at higher risk of SARS-CoV-2 infection (2045 visible minorities and 542 Indigenous 417 individuals). Of these, 1229 agreed to provide DBS and were included in Phase 4 mailouts (919 visible minorities and 310 418 Indigenous individuals). In Phase 5, a subset of 1304 participants who had recently tested negative for antibodies to 419 nucleocapsid (N) were selected for a supplementary DBS sample; in Phase 6, 5703 DBS participants from any previous 420 phase were enrolled.

421 Participants were not compensated financially by the study for participating, but earned modest redeemable points from the

Angus Reid Forum.<sup>29</sup> Figure S1 illustrates the study recruitment and flow; there were few (about 1%) exclusions, mostly from  $\frac{1}{2}$ 

423 incomplete testing.

#### 424 IgG serology

Participants collected five small circles of blood on special bar-coded filter paper, dried the sample for at least two hours, placed it in a two-layer protective pouch, and returned it to St. Michael's Hospital in Toronto, postage prepaid. Mailing time across Canada ranged from about 3 to 8 days. Upon arrival, samples were scanned, catalogued, and stored at 4°C in larger boxes with additional desiccant, and monitored for humidity levels (kept <20%).</p>

Antibodies were then eluted from a 4.7 mm punch in 99  $\mu$ L of PBS + 0.1% Tween (PBS-T) and 1% Triton X-100. The use of 99  $\mu$ L was to ensure sufficient eluate to test three antigens (spike protein, receptor binding domain (RBD) of the spike, and

430 99 µL was to ensure sufficient eluate to test three antigens (spike protein, receptor binding domain (RBD) of the spike, ar 431 nucleocapsid protein (N)). Punches were incubated in elution buffer for a minimum of 4 hours with gentle shaking (150

432 RPM) at room temperature or overnight at 4°C. The samples were then centrifuged at 1000 g for 30 seconds.

433 The Network Biology Collaborative Centre at Sinai Health, Toronto, conducted a high-throughput, highly sensitive 434 chemiluminescence-based ELISA targeting the spike protein, RBD, and N. Chemiluminescent ELISA assays were performed

435 as previously described on a ThermoFisher Scientific F7 robotic platform<sup>13,14</sup> with a few modifications. Briefly, LUMITRAC

- 436 600 high-binding white polystyrene 384-well microplates (Greiner Bio-One #781074, VWR #82051- 268) were pre-coated 437 overnight with 10 µL /well of antigen (50 ng spike (SmT1), 20 ng RBD and 7 ng nucleocapsid, all supplied by the National
- 437 overnight with 10 µL/wen of antigen (50 ng spike (Sin11), 20 ng KBD and 7 ng nucleocapsid, an supplied by the National 438 Research Council of Canada (NRC)). After washing (all washes were 4 times with 100 µL PBS-T), wells were blocked for 1
- 439 hour in 80  $\mu$ L 5% Blocker BLOTTO (ThermoFisher Scientific, #37530) and then washed. 10  $\mu$ L of sample (2.5 or 0.156  $\mu$ L 440 of DBS eluate diluted in 1% final Blocker BLOTTO in PBS-T) was added to each well and incubated for 2 hours at room
- of DBS eluate diluted in 1% final Blocker BLOTTO in PBS-T) was added to each well and incubated for 2 hours at room temperature. After washing, 10  $\mu$ L of a human anti-IgG fused to HRP (IgG#5, supplied by NRC, final of 0.9 ng/well) diluted in 1% final Placebar PL OTTO in PBS T was added to each well followed by a 1 hour in what is a strugger to up and the second second
- in 1% final Blocker BLOTTO in PBS-T was added to each well followed by a 1-hour incubation at room temperature. After 443 4 washes, 10  $\mu$ L of SuperSignal ELISA pico chemiluminescent substrate (diluted 1:4 in MilliQ distilled H<sub>2</sub>0) was added to 444 each well and incubated for 5-8 min at room temperature. Chemiluminescence was read on an EnVision (Perkin Elmer) plate 445 reader at 100 ms/upil using an ultra constitute datastar
- reader at 100 ms/well using an ultra-sensitive detector.
- Each 384-well assay plate included replicates of a standard reference curve of a human anti-spike IgG antibody (VHH72-Fc supplied by NRC)<sup>13</sup> or an anti-nucleocapsid IgG antibody (Genscript, #A02039), positive and negative master mixes of pooled serum samples, human IgG negative control (Sigma, #I4506), and blanks as controls. Negative and/or positive DBS controls (defined using plasma serology results) were included in runs in each phase.
- 450 For each antigen, raw values (counts per second) were normalized to a blank-subtracted point in the linear range of the
- 451 standard reference curve to create a relative ratio (hereinafter referred to as antibody levels). The samples were processed at a
- 452 1:4 dilution of the DBS eluate (2.5  $\mu$ L/well of sample) and 1:64 dilution. We used the former to derive positivity threshold
- 453 and the latter to display antibody level distributions.

#### 454 **Determining positivity**

- 455 There is uncertainty in the measured values of the antibodies to N. We sought to reflect this uncertainty in the confidence
- intervals for prevalence estimates. We used control samples and known positives to estimate the probability of seropositivity
- for each sample, and we used multiple imputation to account for the unknown true seropositivity status. We estimated log relative rates in a model adjusting for age, sex, region, and vaccination status. Using post-stratification, we computed
- estimates and confidence intervals for prevalence in the population and various subgroups, adjusting for the
- 460 representativeness of the sample.
- Figure S2A shows the histogram of logged N-positivity for known laboratory negative control samples within each testing plate (in blue) and antibody levels from known positive samples from phase 4. Known positives are individuals who reported a positive covid-19 test result more than seven days before their DBS was received. We used maximum likelihood estimation to define skew-normal densities for the case and control samples, shown as solid lines. Figures S2B to S2D show histograms
- 465 of observed antibody levels for each phase, along with a fitted density estimated as a mixture of the red and blue densities
- 466 from Figure S2A. We estimated a mixing proportion for each phase (by maximum likelihood), the densities for each
- 467 component are shown in blue and red for the seronegative and seropositive components respectively.
- 468 For each sample, we calculated a probability of seropositivity using Bayes rule. This probability depends on the mixing
- 469 proportion as well as the red and blue densities, as when prevalence is high the threshold should be lowered to reduce false
- 470 negatives. These probabilities are used for multiple imputation, generating 100 datasets where each sample is designated as
- seropositive or seronegative. For grouping subjects as infected and uninfected in the "immunity wall" figures, cutoffs for
- each phase (shown in Figure S2s) are set so that the expected number of false positives and false negatives is identical.
- 473 Prevalence estimates and their confidence intervals use post-stratification, adjusting the study sample to reflect the Canadian
   474 distribution of population by age, sex, region and vaccination status. For each phase, we fit a logistic regression model which
- 474 included vaccination status (no doses v. one or more) and region (British Columbia and Yukon; Prairie provinces, NWT,
- 476 Nunavut; Ontario; Quebec; Atlantic provinces), each of which are interacted with age and sex (and the age-sex interaction).
- 477 We did not interact vaccination status with region, as the number of unvaccinated individuals in the sample was small in
- some regions. We obtained estimates of log relative rates and the accompanying variance matrix for each of the 100 imputed
   datasets and combined them according to Rubin's rule.
- 480 The population by age, sex, province, and vaccination status at each phase are obtained from the Public Health Agency of
- Canada's Infobase.<sup>9</sup> Weights are calculated for each age-sex-region-vaccination group and a weighted average of group-level  $\frac{1}{30}$
- 482 prevalences is computed with standard errors obtained from the delta method.<sup>30</sup>

#### 483 Interferon-Gamma Release Assay (IGRA) T-cell analysis

- We selected a convenience sample of adults in the Ab-C study within urban Toronto. After obtaining consent for re-contact, participants attended either a Unity Health Toronto hospital visit or agreed to a home visit. A phlebotomist collected one tube of venous blood from each participant using 7 mL lithium-heparin blood collection tubes. Blood collection tubes were mixed by inversion, stored at room temperature, and delivered to the St. Joseph's Health Centre laboratory within 16 hours of collection to be refrigerated at 2-8°C.
- 400 collection to be refrigerated at 2-8°C.
- 489 Prior to stimulation, samples were removed from refrigeration for 30 minutes. For each whole-blood sample, one stimulation 490 tube set from the Quan-T-Cell SARS-CoV-2 kit (EUROIMMUN, ET 2606-3003) was warmed to room temperature. Each set 491 consisted of three stimulation tubes: (1) CoV-2 IGRA BLANK: no T-cell stimulation, for determination of the individual 492 IFN-γ background; (2) CoV-2 IGRA TUBE: specific T-cell stimulation using antigens based on the SARS-CoV-2 spike 493 protein; (3) CoV-2 IGRA STIM: unspecific T-cell stimulation by means of a mitogen, for control of the stimulation ability. 494 The blood collection tube was mixed by gentle inversion, then sampled using 1 mL pipets to draw and transfer 500 µL of 495 whole blood to each of the three tubes. The filled stimulation tubes were sealed and mixed by rapid inversion, then shaken by 496 hand and incubated at 37°C for 20-24 hours. At the end of the incubation period, the tubes were removed from the incubator 497 and centrifuged for 10 minutes between 6000-12000 x g.
- 498 Following centrifugation, the plasma obtained from the stimulated whole-blood samples was diluted and used on the anti–
- 499 IFN-γ-coated ELISA plate. EUROIMMUN Mississauga conducted interferon-gamma release assays using the Quan-T-Cell
- 500 ELISA (EQ 6841-9601). 100  $\mu$ L of the calibrators, controls, and diluted plasma samples (1:5 in sample buffer) were
- transferred into the individual microplate wells and incubated for 120 minutes at room temperature. The wells were washed
- 502 (5 times, each using 300  $\mu$ L of wash buffer). 100  $\mu$ L of biotin was pipetted into each well and incubated for 30 minutes at
- 503 room temperature. The wells were washed, and 100  $\mu$ L of enzyme conjugate was pipetted into each well and incubated for 30

504 minutes at room temperature. The wells were washed, and 100 µL of chromogen/substrate solution was pipetted into each

well and incubated for 20 minutes at room temperature, protected from direct sunlight. 100 µL of stop solution was pipetted into each well. Photometric measurements of the colour intensity were made at a wavelength of 450 nm and a reference

507 wavelength between 620 and 650 nm.

#### 508 Epidemiological Analyses

509 This analysis focused on Phases 3 to 6 of the Ab-C study, which correspond to the pre-omicron (Aug 15 to Oct 15, 2021) and 510 omicron (BA.1/1.1, BA.2, and BA.5) periods (Jan 24 to Mar 30; May 27 to Jul 1; and Sep 26 to Nov 21, 2022), respectively. 511 To confirm the Ab-C data is representative of the Canadian population, we calculated the proportion of participants who filled 512 out the survey and provided DBS by demographic characteristics (province, household size, age, sex, education, ethnicity, 513 weight, smoking status, diabetes, hypertension) and vaccination status, and compared these to the Canadian national data 514 (Table S1).

515 As already reported,<sup>4</sup> the demographic and health characteristics of those who completed surveys and provided DBS were 516 generally comparable to the Canadian census population, except for fewer adults with an educational level of some college or 517 less in the Ab-C study compared with the census population. In Phase 6, the proportion of adults unvaccinated was similar in 518 the Ab-C surveyed population (8%) as in Canada overall (10%). However, the unvaccinated rates were lower in those who 519 submitted DBS samples (3%). We have previously found greater unvaccinated rates among the lower levels of education.<sup>12</sup> 520 Education level (some college or less, college graduate, university graduate) was inversely correlated with vaccination status: 521 chi-squared statistic 17.156 (df=2; p-value of 0.0001882). Hence, we adjusted for vaccination status when calculating 522 estimates of cumulative incidence. Moreover, the Ab-C study has had fewer racial or ethnic minority adults (which is defined 523 by Statistics Canada, the national lead statistical agency, as "Visible Minorities") but more Indigenous adults than the census 524 population. Compared with the census population or nationally representative surveys, study participants had a similar 525 prevalence of obesity, current or former smoking, diabetes and hypertension.

526 The phase 3 to 6 population distributions, which are most directly relevant to estimating cumulative and period-specific 527 Omicron incidence, are broadly similar among those who completed surveys and those who provided a DBS (Table S1).<sup>5</sup> 528 Finally, a comparison of those invited who participated and did not in Phase 1 of the study showed a bias towards greater 529 female participation.<sup>4</sup> However, differences by sex were not important predictors of cumulative incidence (data not shown), 530 so this bias does not materially affect the overall estimates of cumulative infection.

531 The age-specific "immunity wall" in Figure 4 defines infection as either having tested positive on polymerase chain reaction 532 or antigen rapid test or with antibodies to the N antigen (which is appropriate among the largely vaccinated cohort). N 533 positivity reflects infection and would not arise from Canadian-approved vaccines that only contain the spike protein. We 534 defined infection as any positive covid-19 test more than 7 days prior to the DBS being received and any N positivity.

535 We obtained the overall cumulative incidence of SARS-CoV-2 infections based on N positivity and derived the 95%

536 confidence intervals using the delta method.<sup>7</sup> In order to examine the level of antibody response from infection and

vaccination (by vaccine doses), we display the distributions of antibodies to spike antigens (at the 1:64 dilution) using box
 plots with jitter (Figure 2). Results for antibodies to RBD are similar (Figure S4). All analyses were performed using Stata 17

539 and R 4.2.1.

#### 540 Appendix 2-figure 1. Study flow including sampling and study inclusion by phase in the Ab-C study







Notes: A) Fitted densities for known cases and controls in calibration samples; B) densities estimated for infected and uninfected individuals from the mixture model in phase 3 samples; C) phase 4 samples; and D) phase 6 samples.

#### 549 Appendix 2-figure 3. Levels of antibodies to RBD stratified by infection and number of vaccination doses.



#### 554 Appendix 2-figure 4. Levels of antibodies to the spike protein stratified by infection, vaccination doses, and time since 555 last vaccination or since last infection.

#### (A) Uninfected by time of last dose



(B) Infected by time of last dose and infection







Notes: X-axis represents interferon-gamma stimulation on blood samples using antigens based on the SARS-CoV-2 spike protein (<u>https://www.coronavirus-diagnostics.com/documents/Indications/Infections/Coronavirus/ET\_2606\_D\_UK\_A.pdf</u>), and y-axis represents SARS-CoV-2 spike protein antibodies in dried-blood spot samples. The two variables had a Spearman correlation of 0.508. Smoothed curves and 95% confidence intervals were obtained using locally weighted scatterplot smoothing with span parameter of 0.8.<sup>27</sup>

#### Appendix 3-table 1. Sample characteristics and representativeness of phases 4 and 6 for online surveys and DBS

samples		-		-			•		
	2016 Canadian	Phase 4 Survey		Phase 4 DBS		Phase 6 Survey		Phase 6	
	Census or		0/	sa	mple		0/	DBS	sample
	national surveys	n 14224	%	n 5021	%	n 10088	%	n 2279	%
		14224	22.60/	5031	24.00/	10088	24 (0/	33/8	22 50/
High risk regions*		4824	33.6%	1/32	34.0%	3530	34.6%	1164	33.5%
Province	290/	5707	40.00/	2102	41 50/	2057	20.20/	1417	41 (0/
	38%	5707	40.0%	2103	41.5%	3957	39.3%	1417	41.0%
British Columbia & Fukon	14%	2802	20.1%	1035	20.9%	2120	21.1%	752	22.4%
Quebec	23%	1/64	12.5%	01/	12.2%	1223	12.0%	364	10.5%
Prairie provinces & NW I	19%	2992	21.1%	979	19.6%	2155	21.5%	055	19.4%
Atlantic provinces	/%	899	6.2%	297	5.9%	633	6.2%	212	6.2%
Sex	400/	(452	45.00/	1007	40.10/	4402	45.00/	1071	27.90/
	49%	6453	45.9%	1997	40.1%	4492	45.0%	12/1	37.8%
Female	51%	/028	55.1%	3003	59.5%	5515	54.2%	2089	01.7%
Prefer to self-describe		143	1.0%	31	0.6%	81	0.8%	18	0.6%
Age group	400/	2(22	22.00/	1000	10.50/	2094	10 10/	510	12.00/
18-39 years	49%	5105	25.9%	1060	19.5%	2084	19.1%	512	15.8%
40-59 years	28%	2202	30.1%	1752	34.4%	3099	35.9%	004	32.7%
60-69 years	12%	3303	24.5%	1355	28.3%	2484	26.1%	994	30.7%
70+ years	11%	2094	15.5%	804	17.9%	1821	18.9%	/38	22.8%
Education	450/	2270	24.00/	1050	20.00	2205	24.20/	702	20.70/
Some college or less	45%	3372	34.0%	1050	30.6%	2395	34.3%	1102	30.7%
College graduate	32%	4680	31.6%	1621	31.5%	3333	32.1%	1102	32.3%
University graduate	23%	6172	34.4%	2360	37.9%	4360	33.6%	1574	37.0%
Visible minority	22%	3438	23.5%	820	15.8%	2482	24.2%	525	15.3%
Indigenous	5%	1504	11.0%	495	10.2%	803	8.6%	234	7.3%
Household size	290/	0(14	10.70/	000	10.70/	2070	20.00/	720	01.50/
Live alone	28%	2614	18.7%	990	19.7%	2070	20.8%	/30	21.5%
Т wo реоріе	34%	6144	44.0%	2328	47.0%	4475	45.2%	1635	49.4%
Three people	15%	2338	16.2%	/4/	14.8%	1558	15.2%	4//	14.0%
Four people or more	22%	3128	21.2%	966	18.5%	1985	18.8%	530	15.2%
Ever smoking $O_{\rm b} = (220  {\rm km/m^2})$	54%	2750	49.8%	1269	49.1%	4/81	50.0%	1599	27.8%
Obesity (≥30 kg/m)	27%	3/50	27.4%	1308	28.1%	2075	27.7%	908	27.8%
Diabetic history	9%	1418	10.6%	518	11.0%	1037	10.9%	359	21.20
Hypertension history	23%	3820	28.4%	1452	30.4%	2850	29.1%	1006	51.5%
vaccination <sup>*</sup>	100/	1075	0.70/	200	1.50/	720	7.00/	05	2 10/
Vaccinated	10%	12/3	9.7%	4910	4.3%	0240	02.10/	2702	0.1%
	90%	12949	90.5%	4819	93.3%	9549 76	92.1%	3283	90.9%
Two doses	1%	2807	20.0%	23	19.5%	1246	12.00/	222	0.5%
1 wo doses	29%	3807	29.9%	8/0	18.3%	1240	15.0%	233	24.00
I nree doses	32%	9006	69.0%	3920	81.0%	3801	57.2%	1158	54.0%
Four or more doses	28%					4226	41.1%	1882	55.5%

\*17 high-burden regions identified from a regression analysis of SARS-CoV-2 case counts

†As of 1 January 2023 

Broadly, the characteristics of the cohort did not change materially between phase 3 and 4 and phase 6, such as in the persistent underrepresentation of lesser educated Canadian adults.<sup>4</sup> 



\* Dotted line indicates period after restrictions to PCR testing eligibility † Major variants include AE.8, B.1.36.36, B.1.2, B.1.160, B.1.438.1

Figure 1. Seven-day rolling averages of PCR-confirmed COVID-19 cases in Canada (black solid and dotted line), and SARS-CoV-2 vaccinations (any dose; red line) in relation to the data collection phases of the Ab-C study. Testing and vaccination data were derived from COVID-19 Tracker Canada as of 3 February 2023 (https://COVID19Tracker.ca).<sup>26</sup> Data on major variants were obtained from Public Health Agency of Canada's Health Infobase COVID-19 epidemiology update (https://health-infobase.canada.ca/covid-19/testing-variants.html).<sup>9</sup> Dotted lines for PCR-based testing after 1 January 2022 reflect the major uncertainty in PCR-based testing. Widespread PCR testing guidelines became stricter and were significantly scaled back in community settings and thus became far less reliable to monitor trends.





Infection only or with 1 vaccine dose n=100

#### Uninfected

2 vaccine doses n=23

3 vaccine doses n=146

4 vaccine doses n=479

2 vaccine doses

#### Infected

n=208

0.0



3 vaccine doses n=1004 4 vaccine doses n=1398

> 0.5 1.0 1.5 2.0 Levels of antibodies to the spike protein (1:64 dilution)

**Figure 2. Levels of antibodies to the spike protein stratified by infection and number of vaccination doses.** Circles represent individuals with their last vaccination (or unvaccinated) >10 days prior to dried blood spot sample collection (n=3378 with complete information available as of the time of analyses after excluding 14 low quality samples). We further excluded 16 participants whose samples were seronegative and viral test was positive, but who did not provide viral test dates or reported test dates less than eight days from the receipt of DBS. The solid-coloured line represents the median and box plots show the interquartile range. The results above a relative level of 1.2 are outside the linear range of the assay. Results using the receptor-binding domain antigen were similar to the spike protein (supplementary figure S3). (A) By age group

(B) By sex



Figure 3. Age, sex, and ethnicity-specific trends to nine months in levels of antibodies to the spike protein among adults vaccinated with 3-4 doses, stratified by infection more than six months ago or less than six months ago. See footnote to figure 2 for testing details. We created smoothed curves and 95% confidence intervals using locally weighted scatterplot smoothing with span parameter of 0.8.<sup>27</sup>





## Figure 4. Cumulative incidence in each stratum of infection and vaccination in the pre-Omicron wave, during the Omicron BA.1/1.1 wave, and during the BA.2 and BA.5 waves by age group.

\*Including uninfected and infected cases. The first column in each age group represents the antibody and viral test positivity for the entire period prior to Omicron, whereas the second column represents the values during the omicron BA.1/1.1 wave and the third during the BA.2/5 waves. By the last time period studied, the numbers of participants aged 15-59 who were N-positive, viral test–positive, and positive to both were 675 (41%), 37 (2%), and 699 (43%). The comparable numbers for participants aged 60 or more were 763 (44%), 35 (2%), and 500 (29%).