analyzed (e.g., brain, liver, spleen, kidney, small and large intestine, heart, and eyelids). These results suggest that virus infection by natural transmission between cats, as well as by experimental inoculation, induces protective immunity against a second SARS-CoV-2 infection.

In conclusion, SARS-CoV-2 replicated effectively in the upper respiratory tract in cats, and infectious virus was cleared from the lungs within 6 days of infection; however, histopathologic examination demonstrated chronic lung sequelae in cats even a month after viral clearance. After initial infection with SARS-CoV-2, cats were protected from reinfection, with no virus replication in respiratory organs and no additional lung damage.

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Long-Term Humoral Immune Response in Persons with Asymptomatic or Mild SARS-CoV-2 Infection, Vietnam

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Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is the causative agent of the coronavirus disease (COVID-19) pandemic (1). Effective vaccines are vital for mitigating the impact of the pandemic. As such, synthesizing a long-term humoral immune response to SARS-CoV-2 remains essential to developing and implementing a SARS-CoV-2 vaccine. We report a longitudinal study of 11 persons with SARS-CoV-2 infection in Vietnam, in which we monitored antibody responses for up to 30 weeks after infection.

We included patients with a confirmed SARS-CoV-2 infection admitted to a COVID-19 treatment center in central Vietnam during January–March 2020. To enable long-term follow-up, we excluded all short-term visitors. We collected information from each participant about clinical status, travel history, contacts with persons with confirmed cases, and personal demographics. For plasma collection, we applied a flexible sampling schedule encompassing 30 weeks after diagnosis, stratified by collection at 1, 2–3, 4–7, and ≥18 weeks after diagnosis.

We measured antibodies against 2 main immunogens of SARS-CoV-2, the nucleocapsid (N) and spike (S) proteins, by using 2 well-validated sensitive and specific serologic assays, Elecsys Anti–SARS-CoV-2 assay (Roche, https://diagnostics.roche.com) (2) and SARS-CoV-2 Surrogate Virus Neutralization Test (sVNT) (GenScript, https://www.genscript.com) (3). The former is an electrochemiluminescence immunosassay that uses recombinant N protein for qualitative detection of pan Ig, including IgG, against SARS-CoV-2. The latter is a surrogate assay for measuring receptor-binding domain–targeting neutralizing antibodies (RBD-targeting NAbs) (3,4), in principle a blocking ELISA that quantifies antibodies that block the receptor–RBD interaction (3). Our study forms part of the national COVID-19 response and was approved by the institutional review board of the Pasteur Institute in Nha Trang, Vietnam.

During the study period, there were a total of 23 patients with confirmed SARS-CoV-2 infection in central Vietnam. Ten were tourists and were thus excluded from the study. Of the remaining 13, a total of 11 consented to participate in this study. Among study participants, 6 were female and 5 were male; the age range was 12–64 years (Table). Seven experienced mildly symptomatic infection and did not require supplemental oxygen during hospitalization; 4 were asymptomatic. Before becoming ill, 3 had traveled to a SARS-CoV-2–endemic country, including patients 2 and 3, who had traveled to Malaysia and patient 4 had traveled to the United States. Patient 4 transmitted the virus to 6 of her contacts, including 4 family members and 2 employees. Of these, 2 transmitted the virus to another family member (Table; Appendix Figure, https://wwwnc.cdc.gov/EID/article/27/2/20-4226-App1.pdf).

### Table. Demographics, travel history, contact history, clinical status, and outcome for participants in study of long-term humoral immune response in persons with asymptomatic or mild SARS-CoV-2 infection, Vietnam, 2020*

| Patient no. | Age, y/sex | Province | Presumed exposure | Symptoms developed | Diagnosed | Presumed incubation period, d | Recent travel history | Contact with confirmed patient | Clinical status | Hospital stay, d |
|-------------|------------|----------|------------------|-------------------|-----------|-----------------------------|----------------------|-----------------------------|-----------------|-----------------|
| 1           | 25/F       | Khanh Hoa | Jan 14           | Jan 18            | Jan 24    | 4                           | None                 | 1 of first 2 cases in Vietnam | Sympt           | 11              |
| 2           | 42/M       | Ninh Thuan | Feb 27–Mar 4     | Mar 9             | Mar 16    | 5–14                        | Malaysia             | Unknown                     | Sympt           | 16              |
| 3           | 36/M       | Ninh Thuan | Feb 27–Mar 4     | Mar 13            | Mar 17    | 9–15                        | Malaysia             | Unknown                     | Sympt           | 15              |
| 4           | 51/F       | Binh Thuan | Feb 22–29        | Mar 5             | Mar 9     | 7–14                        | USA                  | Unknown                     | Sympt           | 25              |
| 5†          | 51/M       | Binh Thuan | Mar 2–9          | Mar 11            | Mar 11    | 2–9                         | None                 | Husband of patient 4         | Sympt           | 23              |
| 6†          | 64/F       | Binh Thuan | Mar 2–10         | Asympt            | Mar 10    | 5–8                         | None                 | Domestic worker of patient 4 | Asympt           | 31              |
| 7†          | 28/F       | Binh Thuan | Mar 7            | Asympt            | Mar 10    | 3                           | Daughter-in-law of patient 4 | Asympt           | 24              |
| 8†          | 28/M       | Binh Thuan | Mar 2–9          | Mar 11            | Mar 11    | 2–9                         | None                 | Son of patient 4             | Sympt           | 23              |
| 9†          | 47/F       | Binh Thuan | Mar 3–8          | Mar 11            | Mar 11    | 3–8                         | None                 | Mother of patient 7          | Sympt           | 23              |
| 10†         | 37/F       | Binh Thuan | Mar 3–8          | Asympt            | Mar 10    | 2–7                         | None                 | Staff of patient 4           | Asympt           | 24              |
| 11†         | 12/M       | Binh Thuan | Mar 3–8          | Mar 11            | Mar 11    | 2–7                         | None                 | Son of patient 10            | Asympt           | 30              |

*All patients made a full recovery. No patients required oxygen. All patients were of Vietnamese nationality. First enrollment was on January 24, 2020, and last was on March 17, 2020. Last follow-up was on August 13, 2020. Asympt, asymptomatic; sympt, symptomatic.
†Patient numbers match those in Figure 1
‡Patients from a cluster involving 3 household transmission chains (Appendix Figure, https://wwwnc.cdc.gov/EID/article/27/2/20-4226-App1.pdf).
We collected 43 plasma samples from 11 participants within 4 time ranges after diagnosis: <1 week (n = 10), weeks 2–3 (n=11), weeks 4–7 (n=11), and weeks 18–30 (n = 11). During the first week after diagnosis, 1 patient (1/10, 10%) had detectable RBD-targeting NAbs, and none had antibodies against N protein. In subsequent weeks, all (100%) participants tested positive by surrogate virus neutralization. Antibodies against N protein were detected in 10/11 (91%) of the samples collected between the second and third weeks after diagnosis and 11/11 (100%) samples collected at subsequent time points (Figure, panel A).

Previous studies have demonstrated that the inhibition percentage measured by surrogate virus neutralization tests correlates well with neutralizing antibody titers measured by conventional virus neutralization assays or plaque-reduction neutralization tests (3,4). In our study, the inhibition percentage was below the assay cutoff in all but 1 plasma sample taken during the first week after diagnosis and then rapidly increased above the assay cutoff at subsequent time points. At weeks 18–30 after diagnosis, the inhibition percentage declined but remained detectable (Figure, panel B).

We demonstrate that antibodies against 2 main structural proteins (S and N) of SARS-CoV-2 in patients with asymptomatic or mild infections were almost undetectable within the first week after diagnosis. Antibodies rapidly increased in subsequent weeks and peaked around weeks 4–7 before declining during the later phase of infection, consistent with previously reported findings (2,5–7). However, few studies have reported the persistence of long-term humoral immune response to SARS-CoV-2 up to 18–30 weeks after diagnosis (5), especially among mildly symptomatic or asymptomatic infected patients.

The titers of RBD-targeting NAbs, which are well correlated with those of neutralizing antibodies, decayed by weeks 18–30 after infection, suggesting that humoral immunity to SARS-CoV-2 infection may not be long lasting. Because neutralizing antibodies are recognized as a surrogate for protection (7–9), follow-up studies beyond this period are needed to more conclusively determine the durability of these long-term responses and their correlation with protection.

Our collective findings offer insights into the long-term humoral immune response to SARS-CoV-2 infection. The data might have implications for COVID-19 vaccine development and implementation and other public health responses to the COVID-19 pandemic.

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Prevalence and Time Trend of SARS-CoV-2 Infection in Puducherry, India, August–October 2020

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We conducted 3 population-based cross-sectional surveys, at 1-month intervals, to estimate the prevalence and time-trend of severe acute respiratory syndrome coronavirus 2 infection in Puducherry, India. Seropositivity rate increased from 4.9% to 34.5% over 2 months and was 20-fold higher than the number of diagnosed cases of infection.

The magnitude of the ongoing pandemic of coronavirus disease (COVID-19), caused by infection with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), has not been fully assessed because most those infected have no or mild symptoms, and thus do not undergo viral nucleic acid or antigen testing (1–3). Determining the proportion of a population that has had infection at various time points is essential for understanding the dynamics of an epidemic in a particular area.

Puducherry district, population ≈1.25 million, is located in southern India. Its earliest recorded case of COVID-19 was in March 2020; it had 7 total cases by the end of May, 67 by end of June, and 663 by end of July 2020 (4). The district followed national COVID-19 management guidelines, including testing all symptomatic persons and their high-risk contacts.

We conducted 3 community-based serologic surveys for SARS-CoV-2 antibodies in Puducherry at 1-month intervals, i.e., during August 11–16, September 10–16, and October 12–16, 2020 (Figure). Each survey included 900 adults selected using a multistage sampling procedure. In the initial stages, we chose 30 clusters, including 21 of 90 urban wards and 9 of 62 villages, using a probability proportional to size with replacement method; this method replicated the urban-to-rural ratio (70:30) of the district’s population. Thereafter, in each cluster, we chose 30 households by systematic random sampling; we collected blood from 1 adult (≥18 years of age) in each household using a modified Kish method (5,6). The data from these surveys represent the cumulative proportion of

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Appendix

Appendix Figure. Illustration showing the possible transmission chains among persons from a cluster of cases involving members of 3 families. Red circles represent symptomatic patients. Blue circles represent asymptomatic patients. Patients are numbered using the same numbering system presented in the table and figure.