

## References

1. Bellentani S, Tiribelli C. The spectrum of liver disease in the general population: lesson from the Dionysos study. *J Hepatol*. 2001;35:531-7. [https://doi.org/10.1016/S0168-8278\(01\)00151-9](https://doi.org/10.1016/S0168-8278(01)00151-9)
2. Lao Statistics Bureau and UNICEF. Lao social indicator survey II, 2017, survey findings report. 2018 [cited 2021 Nov 2]. <https://www.unicef.org/laos/media/306/file/LSIS2017ENG.pdf>
3. Xaydalasouk K, Sayasinh K, Hübschen JM, Khounvisith V, Keomany S, Muller CP, et al. Age-stratified seroprevalence of vaccine-preventable infectious disease in Saravan, Southern Lao People's Democratic Republic. *Int J Infect Dis*. 2021;107:25-30. <https://doi.org/10.1016/j.ijid.2021.04.033>
4. Black AP, Vilivong K, Nouanthon P, Souvannaso C, Hübschen JM, Muller CP. Serosurveillance of vaccine preventable diseases and hepatitis C in healthcare workers from Lao PDR. *PLoS One*. 2015;10:e0123647. <https://doi.org/10.1371/journal.pone.0123647>
5. Hübschen JM, Jutavijittum P, Thammavong T, Samountry B, Yousukh A, Toriyama K, et al. High genetic diversity including potential new subtypes of hepatitis C virus genotype 6 in Lao People's Democratic Republic. *Clin Microbiol Infect*. 2011;17:E30-4. <https://doi.org/10.1111/j.1469-0691.2011.03665.x>
6. Viet L, Lan NTN, Ty PX, Björkqvist B, Hoel H, Gutteberg T, et al. Prevalence of hepatitis B & hepatitis C virus infections in potential blood donors in rural Vietnam. *Indian J Med Res*. 2012;136:74-81.
7. Riondel A, Huong DT, Michel L, Peries M, Oanh KTH, Khue PM, et al. Towards targeted interventions in low- and middle-income countries: risk profiles of people who inject drugs in Haiphong (Vietnam). *BioMed Res Int*. 2020;2020:8037193. <https://doi.org/10.1155/2020/8037193>
8. Schmutz J. The Ta'Oi language and people. *Mon-Khmer Stud*. 2013;42:i-xiii.
9. Mallory MA, Lucic DX, Sears MT, Cloherty GA, Hillyard DR. Evaluation of the Abbott realtime HCV genotype II RUO (GT II) assay with reference to 5'UTR, core and NS5B sequencing [Erratum in: *J Clin Virol*. 2014;61:625]. *J Clin Virol*. 2014;60:22-6. <https://doi.org/10.1016/j.jcv.2014.02.006>
10. Shin SR, Kim YS, Lim YS, Lee JS, Lee JW, Kim SM, et al. Clinical characteristics and treatment outcome of peginterferon plus ribavirin in patients infected with genotype 6 hepatitis C virus in Korea: a multicenter study. *Gut Liver*. 2017;11:270-5. <https://doi.org/10.5009/gnl16163>

Address for correspondence: Antony P. Black, Vaccine-Preventable Disease Laboratory, Institut Pasteur du Laos, Rue Samsenthai, Ban Kao-gnot, Vientiane, Laos; email: a.black@pasteur.la

## Limited Propagation of SARS-CoV-2 among Children in a Childcare Center, Canada, 2021

Anthony Li, Kieran Moore, Lindsay Bowthorpe, Julie Sousa, T. Hugh Guan

Author affiliations: Kingston, Frontenac, Lennox & Addington Public Health, Kingston, Ontario, Canada (A. Li, K. Moore, L. Bowthorpe, J. Sousa, T.H. Guan); Queen's University, Kingston (A. Li, K. Moore, L. Bowthorpe, T.H. Guan); Ministry of Health of Ontario, Toronto, Ontario, Canada (K. Moore)

DOI: <https://doi.org/10.3201/eid2801.211811>

An outbreak of severe acute respiratory syndrome coronavirus 2 with no definitive source and potential exposure to variants of concern was declared at a childcare center in Ontario, Canada, in March 2021. We developed a robust outbreak management approach to detect, contain, and interrupt this outbreak and limit propagation among children.

On March 1, 2021, an infant enrolled at a childcare center in the Kingston, Frontenac, Lennox, and Addington region in Ontario, Canada tested positive for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2); however, no acquisition source was identified. The next day, another 7 children and staff at the facility tested positive, and an outbreak was declared.

We immediately searched for potential transmission events and deployed a public health inspector and nurse team. The infant had last attended the childcare center >3 days before symptom onset, beyond the 48-hour window for exposure risk according to standard guidance (1). Furthermore, the assessment team identified no travel, occupational, or other contact risks. Out of an abundance of caution, we extended the period of communicability (POC) from 48 to 96 hours, which defined the childcare center as an outbreak setting. We identified staff who had recently traveled to regions with high proportions of SARS-CoV-2 variants of concern (VOC). We were concerned that the increased transmissibility and virulence of a potential VOC outbreak in a childcare center could rapidly spread through the community, given recent studies demonstrating SARS-CoV-2 infection and transmission among children (2,3).

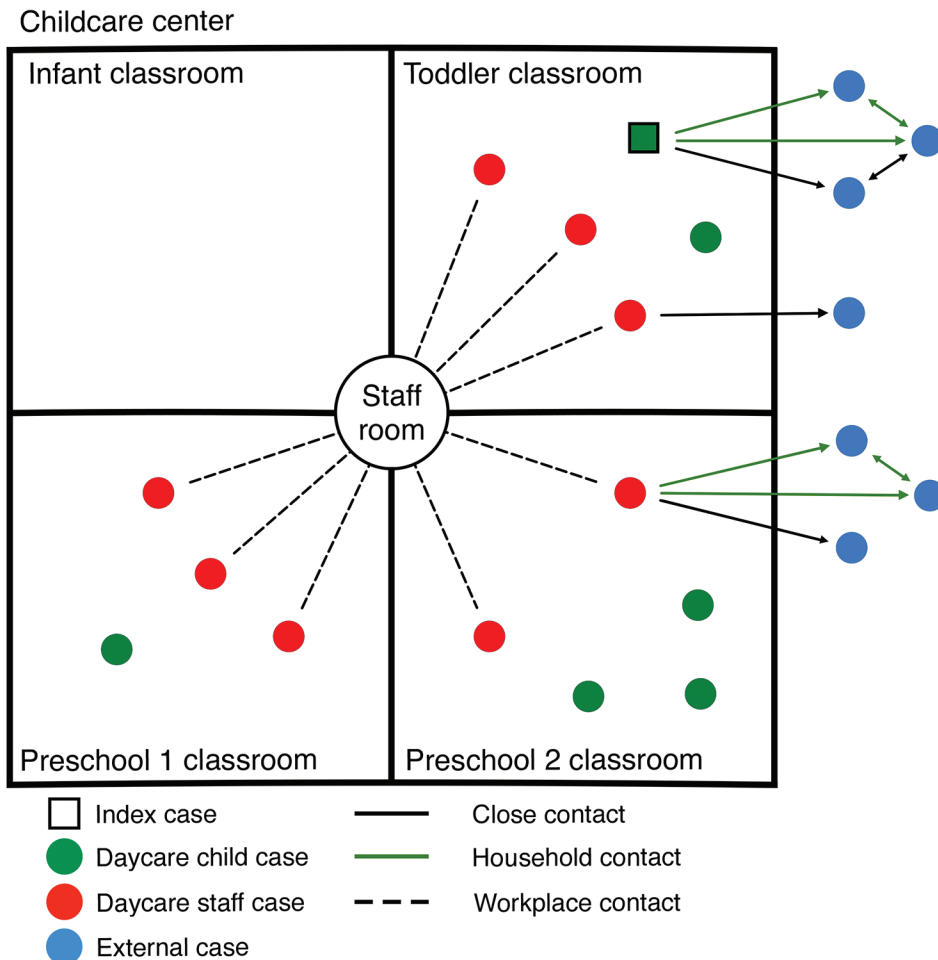
Case investigators gathered symptom profiles, onset dates, detailed exposure histories, risk

factors, and contacts. Because they did not meet early vaccine eligibility criteria, none of the persons had been vaccinated. Because of concern about VOCs, we applied the 96-hour POC to all case-patients and high-risk contacts. Case-patients were required to immediately isolate for 10 days under active monitoring. We advised all high-risk contacts and their household contacts to quarantine for 14 days. As a precaution, we initiated contact tracing before receiving laboratory results for high-risk contacts in whom COVID-19-associated symptoms developed. We requested that all close contacts be tested 3 times while in quarantine: on day 0 and during days 5–7 and 10–12. To be discharged, we required contacts to test negative on days 10–12 or, if having a positive or incomplete test, to quarantine for 10 additional days before retesting. The local Public Health Ontario laboratory conducted real-time reverse transcription PCR testing using the cobas 6800/8800 assay (Roche Molecular Diagnostics, <https://diagnostics.roche.com>) or a

laboratory-developed test (Public Health Ontario, <https://publichealthontario.ca>) (4,5). Testing turn-around time was <24 hours, and positive samples were tested for N501Y and E484K mutations.

A total of 21 SARS-CoV-2 cases were associated with this outbreak during March 1–23, 2021 (Appendix, <https://wwwnc.cdc.gov/EID/article/28/1/21-1811-App1.pdf>): 14 (67%) through direct exposure at daycare and 7 (33%) through secondary transmission (Figure). Average affected age was 22.5 years (range 19 months–68 years); similar proportions of female (11/21) and male (10/21) persons were affected.

For the first generation, the staff attack rate, 47% (8/17), was >4 times higher than the child attack rate, 11% (6/53) (Table) and higher in every classroom with positive cases, aligning with increased SARS-CoV-2 susceptibility and transmission reported among adults compared with children (6–8). Of note, we observed no cases or transmission among nonmobile infants, who remained in assigned cribs in a separate classroom, or their caregivers.



**Figure.** Social network analysis of a COVID-19 outbreak in a childcare center in Ontario, Canada, March 1–23, 2021. The facility had 1 common staff room and 4 physically separated classroom cohorts: infant (6–18 months of age), toddler (18 months–2.5 years of age), and preschool classes 1 and 2 (both 2.5–5 years of age), excluding adult staff.

**Table.** Severe acute respiratory syndrome coronavirus 2 cases and attack rates staff and children during outbreak associated with childcare center, Ontario, Canada, March 1–23, 2021

Category	Infant classroom	Toddler classroom	Preschool 1 classroom	Preschool 2 classroom	Total
<b>Children</b>					
Total no.	6	12	20	15	53
No. cases	0	2	1	3	6
Attack rate, %	0	17	5	20	11
<b>Staff</b>					
Total no.	3	3	8	3	17
No. cases	0	3	3	2	8
Attack rate, %	0	100	38	67	47

We observed proper personal protective equipment, hand hygiene, and cleaning protocols. However, we identified staff breakrooms as high-risk settings because of reduced physical distancing between different staff cohorts and long-term unmasking during meals. Furthermore, some staff were identified as coming to work with COVID-19-associated symptoms which, when combined with high risk for staff exposure, emphasizes the continued importance of careful screening at work and requiring isolation and retesting after a positive test (9). Staff must also be vigilant in adhering to physical distancing and infection prevention and control guidelines (<https://ipac-canada.org>) when socializing outside of the workplace.

Although we identified no definitive acquisition source or transmission incidents, our robust outbreak management approach enabled detection, containment, and interruption of this outbreak with limited propagation among children (Appendix Figure). Extending the POC from 48 to 96 hours broadened our capacity to identify both exposure risks and VOC risk from staff travel. Immediate lockdown of facilities and rapid isolation and quarantine guidance reduced further transmission. The 3-stage testing strategy and short testing turnaround times helped us identify 5 persons who tested positive after initially testing negative (3 on days 5–7 and 2 on days 10–12 of isolation) who might otherwise have further transmitted the virus. Early identification of contacts from second-generation cases and rapid closure, isolation, and testing of other at-risk locations prevented third-generation spread; there was no reported transmission in other workplaces, schools, or the community. We detected no VOCs and presumed this outbreak to involve wild-type strain. No case-patients required hospitalization or died during this outbreak. Our findings show that an aggressive testing protocol, strong collaboration with persons in the outbreak setting, and concentric circle quarantining of contacts were crucial to successfully managing a potential VOC outbreak, particularly when no specific acquisition sources or exposure risks were known.

### Acknowledgments

We thank all the staff at Kingston, Frontenac, Lennox & Addington Public Health who contributed to the investigation of this outbreak, including the investigators, nurses, case and contact management team, and assessment center personnel, as well as the staff at the Kingston Health Sciences Centre and the local Public Health Ontario Laboratory. We also thank Tiffany Ho, Tierra Reay, Carolyn Tran, Brian Mosely, and Adam van Dijk for their contributions to this report. Finally, we acknowledge all the persons and organizations affected by this outbreak and the COVID-19 pandemic.

### About the Author

Anthony Li is a medical student at Queen's University Faculty of Health Sciences, Kingston, Ontario, Canada, who works for Kingston, Frontenac, Lennox & Addington Public Health. His primary research interests include health policy, communicable disease outbreak investigation, and sustainable management.

### References

- Ontario Agency for Health Protection and Promotion (Public Health Ontario). Risk assessment approach for COVID-19 contact tracing. 2021 [cited 2021 Jun 22]. <https://www.publichealthontario.ca/-/media/documents/ncov/main/2020/09/covid-19-contact-tracing-risk-assessment.pdf>
- Paul LA, Daneman N, Schwartz KL, Science M, Brown KA, Whelan M, et al. Association of age and pediatric household transmission of SARS-CoV-2 infection. *JAMA Pediatr.* 2021;175:1151–8. <https://doi.org/10.1001/jamapediatrics.2021.2770>
- Soriano-Arandes A, Gatell A, Serrano P, Biosca M, Campillo F, Capdevila R, et al.; COVID-19 Pediatric Disease in Catalonia Research Group. Household severe acute respiratory syndrome coronavirus 2 transmission and children: a network prospective study. *Clin Infect Dis.* 2021;73:e1261–9. <https://doi.org/10.1093/cid/ciab228>
- Poljak M, Korva M, Knap Gašper N, Fujs Komloš K, Sagadin M, Uršič T, et al. Clinical evaluation of the cobas SARS-CoV-2 test and a diagnostic platform switch during 48 hours in the midst of the COVID-19 pandemic. *J Clin Microbiol.* 2020;58:e00599–20. <https://doi.org/10.1128/JCM.00599-20>
- Corman V, Bleicker T, Brünink S, Drosten C, Landt O, Koopmans M, et al.; World Health Organization. Diagnostic

- detection of 2019-nCoV by real-time RT-PCR. 2020 [cited 2021 Aug 13]. <https://www.who.int/docs/default-source/coronaviruse/protocol-v2-1.pdf>
6. Saheb Sharif-Askari N, Saheb Sharif-Askari F, Alabed M, Temsah M-H, Al Heialy S, Hamid Q, et al. Airways expression of SARS-CoV-2 receptor, ACE2, and TMPRSS2 is lower in children than adults and increases with smoking and COPD. *Mol Ther Methods Clin Dev.* 2020;18:1-6. <https://doi.org/10.1016/j.omtm.2020.05.013>
  7. Mehta NS, Mytton OT, Mullins EWS, Fowler TA, Falconer CL, Murphy OB, et al. SARS-CoV-2 (COVID-19): what do we know about children? A systematic review. *Clin Infect Dis.* 2020;71:2469-79. <https://doi.org/10.1093/cid/ciaa556>
  8. Davies NG, Klepac P, Liu Y, Prem K, Jit M, Eggo RM; CMMID COVID-19 working group. Age-dependent effects in the transmission and control of COVID-19 epidemics. *Nat Med.* 2020;26:1205-11. <https://doi.org/10.1038/s41591-020-0962-9>
  9. Gostic K, Gomez AC, Mummah RO, Kucharski AJ, Lloyd-Smith JO. Estimated effectiveness of symptom and risk screening to prevent the spread of COVID-19. *eLife.* 2020;9:e55570. <https://doi.org/10.7554/eLife.55570>

Address for correspondence: Anthony Li, Kingston, Frontenac, Lennox & Addington Public Health, 221 Portsmouth Avenue, Kingston, ON K7M 1V5, Canada; email: [anthony.li@queensu.ca](mailto:anthony.li@queensu.ca)

## Cluster of SARS-CoV-2 Gamma Variant Infections, Parintins, Brazil, March 2021

Juliana F. da Silva, Roberto J. Esteves, Charlene Siza, Elaine P. Soares, Tatyana C. Ramos, Evelyn C. Campelo, Cristiano F. da Costa, Leila C. de Alencar, Rafaela P. Cavalcante, Clerton R. Florêncio, Tirza P. Mattos, Maria G. Bonecini-Almeida, Luciana Silva-Flannery, Barbara J. Marston, Juliette Morgan, Mateusz Plucinski, Felipe Naveca; CDC Brazil Investigation Team<sup>1</sup>

Author affiliations: US Centers for Disease Control, Atlanta, Georgia, USA (J.F. da Silva, C. Siza, L. Silva-Flannery, B.J. Marston, M. Plucinski); US Centers for Disease Control and Prevention, Brasilia, Brazil (R.J. Esteves, J. Morgan); Secretaria Municipal de Saúde de Parintins, Parintins, Brazil (E.P. Soares,

R.P. Cavalcante, C.R. Florêncio); Fundação de Vigilância em Saúde do Amazonas, Manaus, Brazil (T.C. Ramos, E.C. Campelo, C.F. da Costa, L.C. de Alencar); Laboratório Central de Saúde Pública do Amazonas, Manaus (T.P. Mattos); Fundação Oswaldo Cruz, Rio de Janeiro, Brazil (M.G. Bonecini-Almeida); Instituto Leônidas and Maria Deane, Manaus (F. Naveca)

DOI: <https://doi.org/10.3201/eid2801.211817>

High case counts after the Gamma (P. 1) variant of severe acute respiratory syndrome coronavirus 2 emerged in Brazil raised concerns that previously infected persons might become reinfected. Investigation of a cluster of coronavirus disease cases in Parintins, in the Brazilian Amazon, suggested household transmission but did not identify high rates of reinfection.

In Parintins, Brazil, an increased rate of coronavirus disease (COVID-19)-associated hospitalization, from 75.5 cases/100,000 persons in November 2020 to 397 cases/100,000 persons in February 2021, led to an unprecedented health crisis on this island. The outbreak coincided with emergence of the Gamma (P.1) variant of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), raising concern that the variant was causing infection even in persons who previously had COVID-19 (1). In March 2021, the Municipal Health Department of Parintins, in collaboration with the US Centers for Disease Control and Prevention (CDC), investigated recently infected persons and their household contacts to identify circulating SARS-CoV-2 variants, assess epidemiologic and laboratory evidence of previous SARS-CoV-2 infection in infected persons, and assess intrahousehold transmission.

We used the COVID-19 surveillance database in Parintins to identify persons  $\geq 18$  years of age who had a positive SARS-CoV-2 antigen test result (Panbio COVID-19; Abbott, <https://www.abbott.com>) in the previous 3 days. On March 4 and 5, 2021, the 22 case-patients identified were visited at home, and all adults able to provide written consent were invited to participate; 90 persons (22 index patients, 68 household contacts) agreed. An index case-patient was defined as the person with the earliest symptom onset date in the household; for all but 1 household, index case-patients were the same persons initially identified in the surveillance database. All participants responded to a questionnaire and provided nasopharyngeal swab and dried blood spot samples; nasal swab samples for antigen testing (BINAXNow; Abbott) were obtained from household contacts only.

We tested nasopharyngeal swabs by reverse transcription PCR (RT-PCR) (Allplex 2019-nCoV Assay; Seegene, <https://www.seegene.com>) and by a variant-of-concern-specific RT-PCR protocol (2).

<sup>1</sup>Team members are listed at the end of the article.