#### SUPPLEMENTARY APPENDIX A: METHODOLOGY DETAILS

**Description of Testing Center:** The regional COVID-19 assessment center, termed "Brewer Park", is an assessment center for COVID-19 testing in Ottawa, ON for both adult and pediatric patients. Pediatric patients are seen in a separate section of the center. Tests were booked 0 to 72 hours in advance.

**Study Recruitment Details:** All children presenting to the Assessment Center were asked about their interest in research participation by the daytime Assessment Center staff (separate from the research team). The daytime staff were informed of our research study, and they identified patients who met indications for testing that were similar to our inclusion criteria. Interested participants who were identified by the daytime staff as meeting our inclusion criteria were flagged by the daytime staff, and subsequently approached by our research team. Our research team provided detailed study information to the patients and carried out a full assessment of their eligibility for inclusion. If patients were found to be eligible by the research team was present onsite at almost all hours of the assessment center opening hours, however this varied depending on staffing numbers and availability. The research team provided information on the study and obtained informed consent from all participants.

### List of Variants of Concern in Ontario during Study Period

January 2021: B.1.1.7 (Alpha)

February and March 2021: B.1.351 (Beta)

Inclusion Criteria	*Ottawa Public Health Definition of High risk <sup>1</sup>
1) Children aged 3-17	• A person who lived with someone who has
	tested positive, while that person was not
	self-isolating and infectious (includes
	congregate living settings where direct
	contact (<2 metres) is occurring in shared
	rooms/living spaces
	• A person who provided care for a person
AND AT LEAST ONE OF	who has tested positive (bathing, toileting,
<b>CRITERIA #2-4 BELOW:</b>	dressing, feeding etc.) and/or had direct
2) Child is identified as high risk	contact with infectious bodily fluids (e.g.,
or close contact to a confirmed	coughed on or sneezed on) from up to 2 days
positive case as defined by	(48 hours) before they became sick
Ottawa Public Health*	• A person who had other similar close (<2
	metres) unprotected contact, for more than
AND/OR	15 minutes (the longer they were within 2
3) Travelled outside Canada in the	metres the higher the risk*), from up to 2
past 14 days	days (48 hours) before the person who tested
	positive was sick
AND/OR	

4) Exhibits signs or symptoms of	• Any patient in a healthcare setting in the
COVID-19 viral illness	same room when the person who tested
a. At least one of ("A"	positive was not on droplet and contact
Symptom): Fever,	precautions or other patients in waiting
cough, shortness of	room/common areas (i.e., < 2 metres from
breath, anosmia	person for any duration of time) when the
OR	person who tested positive was not wearing a
b. Two or more of ("B"	surgical/procedure mask
Symptom): Congestion,	• Passengers or crew (e.g. aircraft, train, bus,
sore throat, abdominal	taxi) seated within 2 metres of the person
pain, vomiting, diarrhea,	who has tested positive (approximately two
fatigue, loss of appetite,	seats in all directions) and/or had close
generalized muscle pain,	prolonged contact (more than 15 minutes)
headache	and/or direct contact with infectious bodily
	fluids, while the person who tested positive
	was not wearing a surgical/procedure mask
	• Any person who has travelled outside of
	Canada in the past 14 days

## Table 2: Eligibility Criteria to quality for NP swab PCR test in Canada

Below lists the general eligbility criteria to qualify for a nasopharyngeal swab PCR test in

Canada during the time of our study period.

January 2021	1. Anyone with any symptom consistent with COVID (though kids				
	with minor single sx could watch & wait for 24 hrs)				
	Asymptomatic HRC and those receiving a COVID Alert App				
	<ul> <li>notification</li> <li>Other Asx priority groups: Pre-procedure; Pre-placement into congregate settings; LTCH/RH residents, staff, contractors and visitors; First Nation/Indigenous/Metis/Inuit; International students completing quarantine</li> </ul>				
	4. NOT ELIGIBLE – personal travel, peace of mind, knows a case but				
	no exposure during period of communicability				
February 2021	1. As above, PLUS:				
	2. UK VOC – encourage all returning travellers from UK and				
	their contacts (even if Asx) to be tested				
March 2021	3. As above, PLUS:				
	4. All kids in school/daycare with any symptoms required a				
	test, as their household members were required to isolate				
	pending the results				
	5. Confirmatory testing following + RAT				
	6. LTCH/RH testing was transitioning to RAT programs				
	(several times per week)				

7. Added farm workers and educational workers (teachers,
school bus drivers) to ASx priority group

**Equations:** 

Eq. 1 Kappa =  $\frac{observed agreement-chance agreement}{1-chance agreement}$ 

# Eq. 2 PABAK = 2 \* Observed agreement - 1

Eq. 3:  $Prob((a)symptomatic) = \frac{S \ discordant \ and \ (a)symptomatic}{S(a)symptomatic}$ 

# **SUPPLEMENTARY APPENDIX B: Instruction Manuals for Families**

### Figure 1. Instruction Manual for self-collection saliva kit

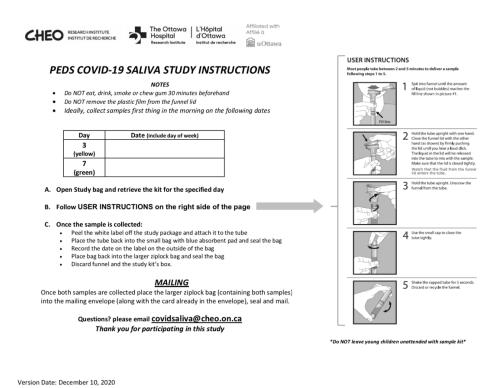
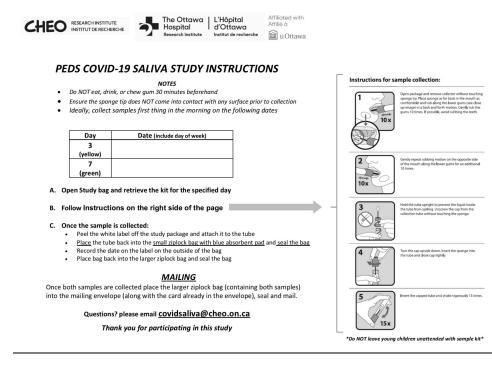


Figure 2. Instruction Manual for self-collection sponge-stick saliva kit



### SUPPLEMENTARY APPENDIX C:

### Part 1: Introduction to longitudinal testing using a simulation

As discussed in the methods, we simulated a testing schedule that lasts for 30 days, with three scenarios where testing is performed every 2, 5, or 7 days, to assess and compare the hypothetical performance of repeat testing using NP- or saliva-based samples. Then we estimated the probability of detecting infection in an individual who was exposed any time before, during or after the testing schedule.

The sequential tests were assumed independent from one another. The simulations were based on modelling the kinetics of the viral load and the analytical sensitivity of the tests (based on NP and saliva samples). The viral load kinetics was modelled as a triangular function on the logarithm scale, parameterized on clinical studies.<sup>2-4</sup> The analytical sensitivity (i.e., sensitivity with respect to viral load present in the sample tested) was modelled with a logistic growth function parameterized on laboratory studies.<sup>5-7</sup> We assumed the analytical sensitivity of the NP-Appendix 1, as supplied by the authors. Appendix to: Hua N, Corsten M, Bello A, et al. Salivary testing for SARS-CoV-2 in the pediatric population: a diagnostic accuracy study. *CMAJ Open* 2022. DOI:10.9778/cmajo.20210279. Copyright © 2022 The Author(s) or their employer(s). To receive this resource in an accessible format, please contact us at cmajgroup@cmaj.ca.

based test to be 98% when the viral load is 10<sup>12</sup> RNA copies per ml, 40% for 10<sup>3</sup> RNA copies/ml and a 99% specificity. For the saliva-based swab, we made the conservative assumption of a lower analytical sensitivity, with 85% when viral load is 10<sup>12</sup> RNA copies per ml (the value of 85% was informed by the estimates reported in the result section of this longitudinal study) and 40% when it is 10<sup>3/0.85</sup> RNA copies per ml. Note this simulation is based on a hypothetical population, not the pediatric cohort reported in this study.

#### Part 2: Model for longitudinal testing

To calculate the overall sensitivity of repeat testing with NP swabs or saliva samples, we modelled the time-dependent logarithmic concentration of SARS-CoV-2 with a triangular function and used a logistic curve to model the analytical sensitivity for both NP and saliva-based assays, making the conservative assumption that the saliva-based assay has a slightly lower analytical sensitivity.

A function, S(v), represents the analytical sensitivity of the assay with respect to a given viral concentration v in the sample tested. The viral load (or concentration) in a patient that was infected a days ago is represented by the function V(a). We model the sensitivity with respect to the time since infection by combining the analytical sensitivity S(v) with the viral kinetics V(a):

$$Se(a) = S(V(a))$$

We model log(V) as a triangular function with units as RNA copies per ml. The shape of this triangular function is informed by clinical studies.<sup>8-10</sup> The triangular function peaks on day 5 after infection at  $10^8$  RNA copies/ml and the limit of detection is reached 13 days after infection (Appendix C Figure 1). Appendix 1, as supplied by the authors. Appendix to: Hua N, Corsten M, Bello A, et al. Salivary

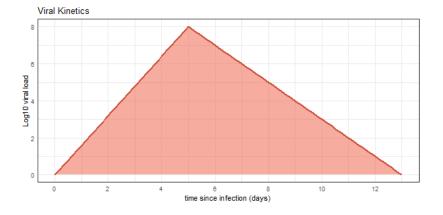
testing for SARS-CoV-2 in the pediatric population: a diagnostic accuracy study. *CMAJ Open* 2022. DOI:10.9778/cmajo.20210279. Copyright © 2022 The Author(s) or their employer(s). To receive this resource in an accessible format, please contact us at cmajgroup@cmaj.ca.

There is a lack of experimental data that would allow to inform the precise shape of *S*. Hence, we model the analytical sensitivity of the test with a logistic function *S* such that S(0) = 0.01 (implicitly represents the test specificity). For the test based on the nasopharyngeal (NP) swab, we assume  $S(10^{12}) = 0.98$  (i.e., the sensitivity of the test is 98% when the viral concentration is  $10^{12}$  RNA copies/ml) and  $S(10^3) = 0.4$ . For the saliva-based test, we assume  $S(10^{12}) = 0.85$  and  $S(10^{3/0.85}) = 0.4$  (Appendix C Figure 1).

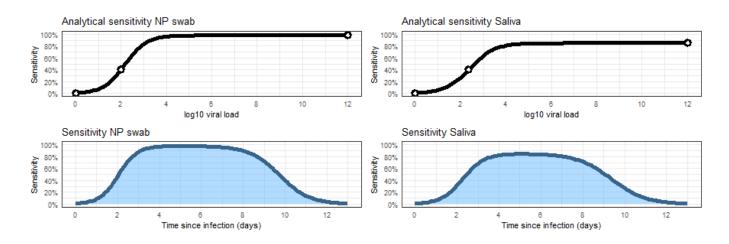
Then, we simulate a schedule of series independent tests (using the same type of assay within the schedule). For an individual infected at time x, and a given testing schedule where the tests occur at times  $t_i$  (i = 1, ..., n), we define the overall sensitivity of the testing schedule as:

$$Se_{overall}(x) = 1 - \prod_{i=1}^{n} (1 - Se(\max(0, t_i - x)))$$

Appendix C Figure 1: Graphical representation of the model assumptions.



Viral Kinetics: Time since infection vs. Log10 Viral Load



Analytical Sensitivity model assumptions

#### Part 3: Interpretation of Longitudinal Testing Results

The following is a description on how to interpret the longitudinal testing result that is described in the main manuscript text, Figure 3.

We simulated a testing schedule that lasts for 30 days, with three scenarios where testing is performed every 2, 5, or 7 days. The x-axis represents the day that an individual is infected, with respect to the day of the first test of the testing schedule (t=0). The y-axis is sensitivity of the test. The vertical dashed lines indicate the day a test is performed.

Using the second graph 'Testing every 5 days' as an example, if an individual gets infected 19 days after the testing schedule starts, they have an 86% chance of testing positive from a saliva sample some time during the testing schedule. This patient will most likely have a negative test on day T=20, because their viral load is not high enough just one day after infection. However, on testing day T=25, their test will likely be positive at that time, because this is 6 days after infection and viral load is likely close to its peak value. Moreover, there is still another chance that on testing day T=30 their test will be (again) positive because infected patients may keep on shedding for a while. Hence, this 86% sensitivity for a patient infected on day T=19 is calculated

by combining all the chances that all the next tests in the testing schedule will be positive: extremely small chance for T=20, large chance for T=25, fair chance for T=30.

As seen in the graphs, the sensitivity drops towards zero at the end of the testing schedule. For a patient infected on day T=29, there is just one test (chance) left on day T=30 to identify their infection, which is very unlikely (just one day of infection is likely not enough to be detected by a test).

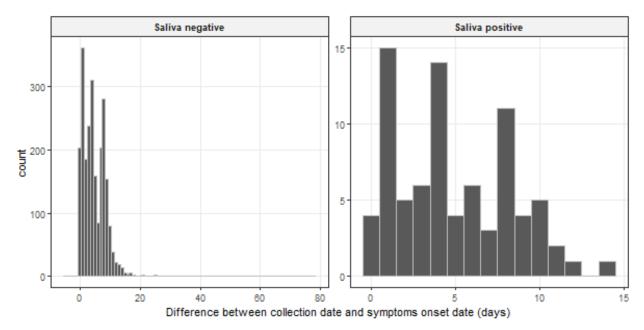
Based on this simulation, a patient that is tested more frequently has a higher chance to be picked up on salivary testing, thereby increasing the sensitivity of the test. This highlights the importance of repetitive testing and highlights the feasibility of using saliva testing in COVID-19 surveillance programs.

## SUPPLEMENTARY APPENDIX D: Additional Results

	Concordant		Discordant		
	Positive	Negative	NP swab positive	Saliva-based assay positive	Total
Symptomatic:	33	994	1	4	1032
Asymptomatic:	11	526	7	4	548
Total	44	1520	8	8	

**Table 1:** Stratification of participants' concordant status by presence or absence of symptoms.

**Figure 1:** Distribution of the difference of time between saliva sample collection date and reported symptom onset date; (A) Distribution by absolute count; (B) Distribution by density *(A)* 



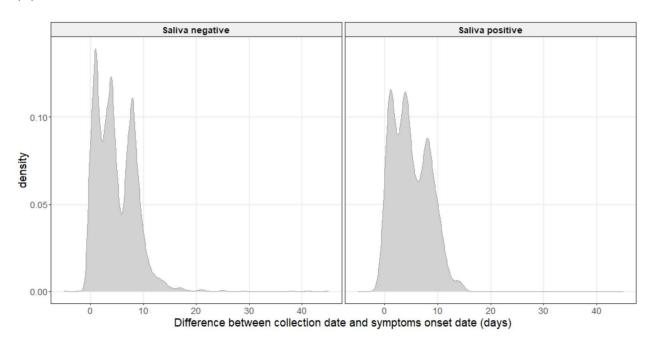
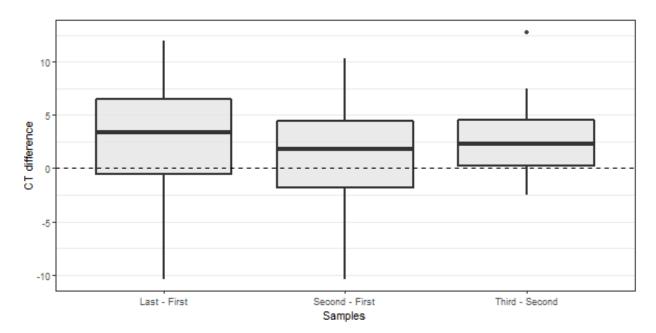


Figure 2: Difference of CT values for saliva samples.



(B)

"Last - First" is the calculation between the first and last sample (irrespective of the total number

of positive samples, as long as there are at least two).

"Second - First" is the calculation between the second sample that was positive and the first that

was positive.

"Third - Second" is the CT difference between the third and second samples that were positive

(only for patients that had three positive results for their saliva-based samples).

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