

1 **Initial evaluation of a mobile SARS-CoV-2 RT-LAMP testing strategy**

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28 **Abstract (220 of 220 words)**

29 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) control in the United States remains hampered, in  
30 part, by testing limitations. We evaluated a simple, outdoor, mobile, colorimetric reverse transcription loop-  
31 mediated isothermal amplification (RT-LAMP) assay workflow where self-collected saliva is tested for SARS-  
32 CoV-2 RNA. From July 16 to November 19, 2020, 4,704 surveillance samples were collected from volunteers and  
33 tested for SARS-CoV-2 at five sites. A total of 21 samples tested positive for SARS-CoV-2 by RT-LAMP; 12 were  
34 confirmed positive by subsequent quantitative reverse-transcription polymerase chain reaction (qRT-PCR) testing,  
35 while eight were negative for SARS-CoV-2 RNA, and one could not be confirmed because the donor did not consent  
36 to further molecular testing. We estimated the false-negative rate of the RT-LAMP assay only from July 16 to  
37 September 17, 2020 by pooling residual heat-inactivated saliva that was unambiguously negative by RT-LAMP  
38 into groups of six or fewer and testing for SARS-CoV-2 RNA by qRT-PCR. We observed a 98.8% concordance  
39 between the RT-LAMP and qRT-PCR assays, with only five of 421 RT-LAMP negative pools (2,493 samples)  
40 testing positive in the more sensitive qRT-PCR assay. Overall, we demonstrate a rapid testing method that can be  
41 implemented outside the traditional laboratory setting by individuals with basic molecular biology skills and can  
42 effectively identify asymptomatic individuals who would not typically meet the criteria for symptom-based testing  
43 modalities.

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55 **Introduction**

56 More than 340,000,000 severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) diagnostic tests have been  
57 performed in the United States as of February 22, 2021, yet it is estimated that 80-95% of infected individuals are  
58 not tested <sup>1,2</sup>. The availability of diagnostic testing for population surveillance around the United States has been  
59 limited because of testing supply shortages and guidelines set by public health officials <sup>3,4</sup>. Multiple studies have  
60 shown that asymptomatic and presymptomatic individuals infected with SARS-CoV-2 can be as infectious as  
61 symptomatic individuals <sup>5-9</sup>, with recent estimates of up to 59% of transmission coming from asymptomatic or  
62 presymptomatic individuals <sup>10</sup>. Virological assessments of SARS-CoV-2-positive individuals and coronavirus  
63 disease 2019 (COVID-19) patients further support the reports of asymptomatic transmission, identifying no  
64 significant differences in viral loads found in the upper respiratory tracts of asymptomatic and symptomatic  
65 individuals <sup>5,7,11-13</sup>. Furthermore, Arons et al. (2020) demonstrated that positive viral cultures can be isolated from  
66 presymptomatic patients up to six days before the onset of symptoms <sup>5</sup>.

67

68 Delays in reporting test results can prevent timely isolation of infected individuals. Since transmission can occur  
69 before symptoms manifest, reporting delays create a major barrier to safely returning to workplaces and schools <sup>14</sup>.  
70 Therefore, there remains an urgent need for rapid tests that identify presymptomatic and asymptomatic individuals  
71 while conserving diagnostic testing reagents. Non-diagnostic point-of-care (POC) testing, used in conjunction with  
72 the current clinical diagnostic testing regimen, may improve our ability to identify infectious individuals and limit  
73 their exposure to others while they are most contagious and conserve clinical diagnostic tests for those who require  
74 confirmatory testing. Incorporating active surveillance using POC tests as part of mitigation strategies for reopening  
75 K-12 schools could play an integral role in reducing SARS-CoV-2 transmission among students, teachers and staff  
76 members, families, and the surrounding community <sup>15,16</sup>.

77

78 Loop-mediated isothermal amplification (LAMP) is a low-cost method for rapid target-specific detection of nucleic  
79 acids <sup>17</sup>. LAMP has long been used as an alternative to gold-standard quantitative reverse transcription polymerase  
80 chain reaction (qRT-PCR) to surveil populations for a variety of pathogens, especially in resource-limited settings  
81 <sup>18-22</sup>. Reverse transcription LAMP (RT-LAMP) assays have recently been developed for rapid SARS-CoV-2 testing

82 <sup>23–29</sup>. RT-LAMP is an appealing candidate for POC SARS-CoV-2 testing because it is inexpensive, circumvents  
83 supply shortages by relying on different reagents than current diagnostic tests, requires minimal sample processing,  
84 and can be deployed outside of traditional laboratory settings. Recently, a number of studies have shown the  
85 correlation between the presence of virus in saliva and nasopharyngeal swabs, demonstrating that saliva specimens  
86 are a valid and reliable alternative to nasopharyngeal swab specimens for SARS-CoV-2 testing <sup>30–35</sup>. Saliva  
87 specimen self-collection is noninvasive, can be done at home, does not require swabs or personal protective  
88 equipment, and limits direct contact between test operators and testing populations. Here we describe our experience  
89 implementing a simple, rapid-turnaround, mobile, non-diagnostic SARS-CoV-2 testing workflow combining self-  
90 collected saliva and RT-LAMP in volunteers without symptoms of SARS-CoV-2 infection. Individuals were  
91 strongly encouraged to isolate and obtain follow-up diagnostic testing after receiving a positive result by RT-LAMP.  
92 This addresses a key knowledge gap of how on-site RT-LAMP testing performs in real-world conditions, since  
93 virtually all previous studies have only evaluated SARS-CoV-2 RT-LAMP in well-equipped molecular biology  
94 laboratories.

95

## 96 **Materials and Methods**

### 97 POC testing sites

98 To begin operating voluntary POC testing, we developed a system of color-coded storage bins for equipment and  
99 supplies, as well as assembled folding tables, chairs, extension cords, and coolers that could be easily  
100 decontaminated and packed to fit in a Dodge Caravan (FCA US LLC., Auburn Hills, MI) or other, similarly sized  
101 minivan for transportation between testing sites and our base laboratory facility. On July 16, 2020, we launched our  
102 first mobile POC testing sites which ultimately expanded over 18 weeks to include two workplaces, two K-12  
103 schools, and an athletics program (Suppl. Table 1). With the exception of the athletics program, sites were initially  
104 outdoors, sometimes under an overhang, but otherwise open to the environment. The athletics site was a climate-  
105 controlled, indoor practice field. At all sites, equipment and reagents were transported by minivan and surfaces were  
106 disinfected during assembly, breakdown, and frequently throughout testing. Participant consenting and volunteer  
107 sample collection were performed on-site but separated from the sample preparation and assay areas (most  
108 commonly on the other side of the building). In an effort to limit contamination, each assay area was set up with

109 three separate folding tables: (1) sample heat-inactivation and preparation, (2) preparation of RT-LAMP reagents  
110 and assay set-up, and (3) RT-LAMP incubation and imaging. Individuals responsible for sample inactivation and  
111 performing assays wore appropriate personal protective equipment (PPE) including N95 face masks, face shields  
112 or safety glasses, disposable lab coats, and double gloves. In anticipation of wet and cold fall weather, by September  
113 2020, assay workspaces were transitioned to biosafety hoods in a vacant indoor laboratory space for several POC  
114 testing locations. In October 2020, we received IRB approval for obtaining consent for repeat SARS-CoV-2 testing.  
115 This allowed us to transition away from consenting participants at each testing time point and instead allowed each  
116 enrolled participant to consent once regardless of the number of times they supplied a sample. Following reports  
117 that SARS-CoV-2 RNA is stable in saliva at room temperature for prolonged periods <sup>36</sup>, we also transitioned away  
118 from in-person sample collection at some of the testing sites and instead distributed self-collection take-home kits  
119 for drop off at designated locations for same day processing.

120

#### 121 Sample collection and preparation

122 We obtained approval from the University of Wisconsin-Madison Institutional Review Board (#2020-0855 and  
123 #2020-1142). Participants were advised to avoid eating or drinking anything except for water for 30 minutes prior  
124 to providing a sample. After providing informed consent, volunteers self-collected at least 50  $\mu$ l of saliva in a 1.5  
125 ml “safe-lock” microcentrifuge tube using a 1000  $\mu$ l unfiltered pipette tip to funnel the specimen into the tube. Each  
126 volunteer disinfected the outside of the tube with a pre-moistened disinfectant wipe. Samples collected in-person  
127 were typically processed within three hours of collection through our RT-LAMP mobile testing workflow, while  
128 samples collected using take-home kits were typically processed within 30 hours (Figure 1). Samples were first  
129 incubated in a heat block at 65°C for 30 minutes to inactivate SARS-CoV-2 <sup>37</sup> and then incubated in another preset  
130 heat block at 98°C for three minutes to improve nucleic acid detection and inactivate salivary enzymes <sup>38</sup>. The  
131 inactivated saliva was then centrifuged for two minutes in a benchtop microcentrifuge. Fifty microliters of the saliva  
132 supernatant were then added to 50  $\mu$ l of 1x phosphate buffered saline, pH 7.4 (1x PBS).

133

#### 134 RT-LAMP reactions

135 Three microliters of the saliva/PBS mixture for each sample were added in duplicate to 17  $\mu$ l of a colorimetric RT-

136 LAMP reaction mix containing WarmStart colorimetric LAMP mastermix (NEB, catalogue# M1800), water, and a  
137 set of six SARS-CoV-2-specific RT-LAMP primers designed against the N gene<sup>38</sup>. The SARS-CoV-2 RT-LAMP  
138 primer set was previously designed by Broughton et al. and is currently used in an FDA emergency use authorized  
139 (EUA) COVID-19 test by Color Genomics (Table 1)<sup>39, 40</sup>. Reactions were incubated for 30 minutes at 65°C. A  
140 smartphone or tablet was used to record images of each reaction before (time = 0) and after the incubation period  
141 (time = 30). A color change from pink/orange to yellow in at least one of two replicates was scored relative to  
142 gamma-irradiated SARS-CoV-2 (irSARS-CoV-2, BEI Resources, Manassas, VA) that was directly added to RT-  
143 LAMP reactions as a positive control in each batch of reactions at concentrations ranging from 220-3,333 copies/ $\mu$ l  
144 ( $2.2 \times 10^5$  -  $3.33 \times 10^6$  copies/ml). irSARS-CoV-2 was diluted and aliquoted as ready-to-run positive control standards  
145 and stored at -80°C. On the day of testing, the positive controls were removed from the freezer and stored on ice at  
146 POC sites. Individuals whose samples were recorded as potentially positive for SARS-CoV-2 by RT-LAMP were  
147 contacted by an infectious disease clinician in accordance with the IRB protocol and urged to obtain a clinical  
148 diagnostic test to confirm findings and self-isolate in accordance with public health recommendations.

149

150 Limit of detection (LOD) estimation using contrived saliva samples

151 To estimate the limit of detection of the RT-LAMP assay, contrived positive saliva samples were prepared by adding  
152 irSARS-CoV-2 initially diluted in nuclease-free water directly into unaltered saliva collected from a total of 20  
153 SARS-CoV-2-negative individuals with the final dilutions ranging from  $1 \times 10^4$ - $10$  copies/ $\mu$ l ( $1 \times 10^7$ -  
154  $1 \times 10^4$  copies/ml). Dilutions were based on independent, in-house qRT-PCR experiments showing that the ir-SARS-  
155 CoV-2 stock concentration  $8.79 \times 10^6$  copies/ $\mu$ l ( $8.79 \times 10^9$  copies/ml). Seven dilutions of irSARS-CoV-2 were  
156 prepared for each saliva sample in duplicate. RT-LAMP reactions were set up as described previously. Negative  
157 controls consisting of saliva from each of the donors without addition of irSARS-CoV-2 were also prepared in  
158 duplicate. Reactions were called positive if a color change from pre-amplification to post-amplification occurred in  
159 at least one of two replicates that was consistent with that of positive controls (a clean yellow color).

160

161 Limit of detection (LOD) estimation using clinical samples

162 De-identified discard saliva samples from 38 SARS-CoV-2-positive patients were provided by the University of

163 Wisconsin Hospitals and Clinics (UWHC) for evaluation of RT-LAMP performance with known positive saliva  
164 samples. Clinical saliva samples were originally collected and stored at 4°C for up to four weeks prior to assessment  
165 by RT-LAMP. Additional 10-fold and 100-fold dilutions were prepared for 13 of the positive clinical saliva samples  
166 in additional saliva collected from a negative volunteer. Clinical samples and dilutions of 13 of those samples were  
167 prepared as described previously except that 20-50 µl of heat-inactivated sample, dependent on total sample volume,  
168 was added to an equal volume of 1x PBS in a clean 1.5 ml screw-top tube and pipetted gently to mix. For each  
169 sample, three microliters were then added to duplicate colorimetric RT-LAMP reactions. Negative and positive  
170 control reactions (described previously) were also prepared in duplicate except that saliva collected from a negative  
171 volunteer was used as the negative control for these reactions. RT-LAMP reactions were prepared and images  
172 collected as described previously.

173

#### 174 Quantitative RT-PCR

##### 175 POC samples

176 We measured vRNA concentration using sensitive qRT-PCR in a subset of the inactivated saliva samples described  
177 above after initial evaluation using RT-LAMP. From July 16 until September 17, saliva samples that were negative  
178 for SARS-CoV-2 by RT-LAMP were pooled into groups of six or fewer for qRT-PCR to balance cost effectiveness  
179 with reasonable estimated detection sensitivity. Ten additional, individual RT-LAMP-negative samples were  
180 submitted as negative controls alongside samples identified as positive by RT-LAMP. Saliva samples that were  
181 identified as positive for SARS-CoV-2 by RT-LAMP were tested by qRT-PCR individually to estimate our POC  
182 LOD. RNA was isolated from up to 150 µl saliva and combined with an equivalent volume of nuclease-free water  
183 using the Viral Total Nucleic Acid kit for the Maxwell RSC instrument (Promega, Madison, WI) following the  
184 manufacturer's instructions. Viral load quantification was performed using a sensitive qRT-PCR assay developed  
185 by the CDC to detect SARS-CoV-2 (specifically the N1 assay) and commercially available from IDT (Coralville,  
186 IA). The assay was run on a LightCycler 96 or LC480 instrument (Roche, Indianapolis, IN) using the Taqman Fast  
187 Virus 1-step Master Mix enzyme (Thermo Fisher, Waltham, MA). The limit of detection of this assay is estimated  
188 to be 0.2 genome equivalents/µl (200 genome equivalents/ml) saliva. To determine the vRNA load, samples were

189 interpolated onto a standard curve consisting of serial 10-fold dilutions of *in vitro* transcribed SARS-CoV-2 N gene  
190 RNA kindly provided by Nathan Grubaugh (Yale University) and described by Dudley et al. <sup>35</sup>.

191

192 Clinical samples

193 Quantitative RT-PCR was performed using the conditions described above for each of the 38 SARS-CoV-2 positive  
194 saliva samples individually; however, sample volume limitations required that for some samples, only 100 µl saliva  
195 was combined with 100 µl of nuclease-free water prior to RNA isolation. In addition, sample UWHC3 contained a  
196 lower volume than the remaining 37 samples so 50 µl saliva was combined with 50 µl nuclease-free water and used  
197 for RNA isolation as described previously. Viral loads in copies per microliter and corresponding cycle threshold  
198 numbers (Ct) are reported in Table 2.

199

## 200 Results

201 LOD estimation using contrived saliva samples

202 We assessed the LOD for minimally processed saliva samples collected from 20 volunteers by RT-LAMP using  
203 irSARS-CoV-2 spiked into negative saliva samples (Figure 2D). We detected irSARS-CoV-2 by RT-LAMP in two  
204 of two replicates (Figure 2A) at  $2.5 \times 10^3$  copies/µl ( $2.5 \times 10^6$  copies/ml) for 100% of samples, at  $1 \times 10^3$  copies/µl  
205 ( $1 \times 10^6$  copies/ml) for 47.4% of samples, and at 500 copies/µl ( $5 \times 10^5$  copies/ml) for 26% of samples. When we  
206 included samples called positive in at least one of two replicates (see Methods and Figure 2B), the percentage of  
207 contrived samples positive by RT-LAMP at each of the aforementioned dilutions were 100%, 89.5%, and 53%  
208 respectively (Figure 2B). One sample was omitted from the analysis because it turned yellow-orange at all dilutions  
209 before the RT-LAMP reaction incubation began and was therefore uninterpretable. Because in POC testing we  
210 defined a positive RT-LAMP result as an observed post-incubation color change to yellow in at least one replicate,  
211 these results suggested that our 90% LOD is approximately  $1 \times 10^3$  copies/µl ( $1 \times 10^6$  copies/ml).

212

213 LOD estimation using clinical samples

214 To assess the performance of SARS-CoV-2 RT-LAMP in known SARS-CoV-2 positive saliva samples as opposed  
215 to contrived positive samples, we acquired deidentified, discarded saliva samples collected from 38 patients with



216 laboratory confirmed SARS-CoV-2 from UWHC. Nineteen of 38 undiluted qRT-PCR-confirmed positive saliva  
217 samples were also positive for SARS-CoV-2 in two of two replicates by RT-LAMP (Figure 3; Table 2). Two  
218 additional samples were positive in one of two replicates. Quantitative RT-PCR data showed that the viral RNA  
219 (vRNA) loads of the positive samples ranged from 131 copies/ $\mu$ l to  $5.7 \times 10^4$  copies/ $\mu$ l ( $1.31 \times 10^5$ - $5.71 \times 10^7$  copies/ml)  
220 which was consistent with our LOD for contrived samples (Table 3). Positive clinical saliva samples that were  
221 negative by RT-LAMP had estimated vRNA loads ranging from 0.402- $5.49 \times 10^4$  copies/ $\mu$ l. All of the samples that  
222 were negative by RT-LAMP, with the exception of UWHC34 ( $5.49 \times 10^4$  copies/ $\mu$ l), had vRNA loads below our  
223 estimated reliable LOD. Furthermore, for the 13 positive clinical saliva samples that were diluted 10-fold and 100-  
224 fold in additional saliva collected from a negative volunteer, detection decreased with increasing dilution factor  
225 (Table 4).

226

#### 227 POC SARS-CoV-2 RT-LAMP testing

228 From July 16 to November 19, 2020, SARS-CoV-2 RT-LAMP was used to test a total of 4,704 samples collected  
229 from five locations. Participants were enrolled into the study regardless of their SARS-CoV-2 symptom status on  
230 the day of testing. Seventy-one percent of the samples were obtained from individuals at two research facilities,  
231 11% from two K-12 schools, and 18% from an athletics program (Supplemental Table 1). A total of 21 samples  
232 were identified as positive for SARS-CoV-2 by RT-LAMP based on a colorimetric change from pink/orange to  
233 yellow in at least one of two sample replicates (see Figure 2B for example). Similar to our experience with our  
234 contrived LOD samples, about 0.40% (19/4,704) of samples collected during POC testing exhibited a color change  
235 to yellow prior to RT-LAMP assay amplification and were therefore uninterpretable. Follow up qRT-PCR testing  
236 was conducted on each sample that appeared positive after the 30-minute amplification reaction throughout the  
237 study to determine vRNA load. Twelve of the 21 samples called positive in RT-LAMP had detectable SARS-CoV-  
238 2 RNA by qRT-PCR. Viral RNA loads of these samples ranged from 8.58 copies/ $\mu$ l to  $3.62 \times 10^5$  copies/ $\mu$ l ( $8.58 \times 10^3$   
239 copies/ml- $3.62 \times 10^8$  copies/ml) with a median of 504.5 copies/ $\mu$ l ( $5.04 \times 10^5$  copies/ml) (Table 4). Eight of the saliva  
240 samples identified as positive by RT-LAMP were negative by qRT-PCR, suggesting that they were false-positive  
241 RT-LAMP results (approximately 40% of samples called positive by RT-LAMP, 0.17% of total samples tested).  
242 One RT-LAMP-positive sample was not tested by qRT-PCR because the participant did not consent to additional

243 molecular testing. For volunteers who consented to additional research testing from July 16 to September 17, qRT-  
244 PCR testing was conducted for pools of six or fewer for all residual, heat-inactivated samples that appeared  
245 unambiguously negative by RT-LAMP. A total of 421 RT-LAMP-negative pools (2,493 samples) were tested to  
246 estimate the number of SARS-CoV-2-positive samples missed by RT-LAMP. Quantitative RT-PCR detected  
247 SARS-CoV-2 nucleic acids in five pools of RT-LAMP-negative samples. Four out of five of the positive pools  
248 contained levels of SARS-CoV-2 that were below the estimated LOD range for RT-LAMP using crude samples  
249 with vRNA load estimates of 0.236, 0.444, 0.460, 37.5, and 142 copies/ $\mu$ l (236, 444, 460,  $3.75 \times 10^4$ , and  $1.42 \times 10^5$   
250 copies/ml). Taken together, the low prevalence of SARS-CoV-2 in our volunteer testing population (0.36%,  
251 including RT-LAMP-negative, qRT-PCR-positive pools) and the low vRNA load of pools positive by follow-up  
252 qRT-PCR, suggest that these five pools likely contained only a single positive sample each and suggests a false-  
253 negative rate of 0.02% (5/2,493 pools) (Table 4).

254

## 255 **Discussion**

256 Strategic surveillance testing of asymptomatic individuals has been suggested as an important mitigation strategy  
257 for places at high risk for close contact, indoor SARS-CoV-2 transmission: schools, workplaces, places of worship,  
258 and prisons, among others. Decentralized, mobile RT-LAMP-based POC testing workflows can provide same-day  
259 results which can enable people with potential SARS-CoV-2 infections to quickly self-isolate and then obtain  
260 confirmatory diagnostic testing. The low per-test cost (approximately \$7 per sample tested in duplicate) allows for  
261 repeated testing to identify incident infections and reduce the duration of a potentially infected individual's exposure  
262 to others. While RT-LAMP is not as sensitive as diagnostic qRT-PCR tests in laboratory testing, qRT-PCR tests  
263 require centralized labs, which in turn leads to lengthy turnaround times. Over a period of 18 weeks, we performed  
264 4,704 SARS-CoV-2 tests across five sites using a simple, saliva-based, direct RT-LAMP assay. This work  
265 demonstrates the scalability of decentralized, mobile RT-LAMP-based testing and addresses a key knowledge gap  
266 of how POC RT-LAMP testing performs outside of well-equipped molecular biology laboratories.

267

268 Our experiment using direct RT-LAMP with contrived saliva samples from a total of 20 donors demonstrated an  
269 approximate LOD of  $1 \times 10^3$  copies/ $\mu$ l (89.5% in at least one replicate). Overall, our data suggest that the actual LOD

270 for RT-LAMP without RNA isolation may be dependent on the individual sample due to heterogeneity of saliva  
271 pH and composition<sup>41-43</sup>. The RT-LAMP results for 38 clinical saliva samples obtained from SARS-CoV-2-positive  
272 individuals at the UWHC, were consistent with those for the contrived samples. We recognize that more clinical  
273 samples are required for a comprehensive clinical validation, but the LOD observed in clinical samples is further  
274 supported by the low vRNA loads obtained from qRT-PCR-confirmed SARS-CoV-2-positive samples identified in  
275 our volunteer population (Table 4). The performance of our RT-LAMP POC testing workflow demonstrates that  
276 inexpensive, mobile testing can be successfully performed outdoors or in other non-traditional laboratory settings  
277 to identify SARS-CoV-2-positive individuals regardless of whether or not symptoms are present. Our observed  
278 SARS-CoV-2 RT-LAMP positivity rate was 0.25% (12/4,704) for samples confirmed by follow-up qRT-PCR.  
279 Interestingly, the positivity rate of 0.25% in our volunteer population was lower than expected given the disease  
280 activity in our region during this period of time was listed as “critically high”, particularly between September 1  
281 and November 19, 2020 when the county had a 5.42% positivity rate (19,031 positive tests out of 350,722)<sup>44, 45</sup>.  
282 The low positivity rate in our volunteer population may be partly explained by the fact that 71% of tested saliva  
283 specimens came from two research facilities where mask wearing and physical distancing guidelines were  
284 implemented early in the pandemic and followed relatively stringently (Supplemental Table 1). Volunteers for  
285 nonsymptomatic research testing might also have a different risk profile from the overall population.

286

287 Potential drawbacks of colorimetric RT-LAMP-based surveillance for SARS-CoV-2 as described here include the  
288 fact that minimally-processed saliva can result in variable reaction color change without the presence of the target  
289 RNA. However, modifications of RT-LAMP-based SARS-CoV-2 assays to reduce saliva sample variability,  
290 improve result ambiguity, and increase throughput have recently been reported elsewhere and may improve the  
291 implementation of RT-LAMP-based assays for POC use<sup>46-50</sup>. In addition, we relied on a manual RT-LAMP format  
292 during POC testing. Reading assays “by eye” inevitably results in a somewhat subjective determination of positives.  
293 Reducing false-positive results in our POC volunteer populations required consistent use of duplicate reactions for  
294 each individual, which reduced assay throughput and increased the per-sample cost. Even with our efforts to reduce  
295 calling false positive results in our volunteer populations, we still were unable to confirm approximately 40% of  
296 RT-LAMP-positive samples by follow-up qRT-PCR. Whether these false positives resulted from the individual

297 sample variability across saliva donors or temporary storage of the samples prior to follow-up qRT-PCR is unclear  
298 but because volunteers with a potential positive finding were strongly encouraged to receive follow-up,  
299 confirmatory diagnostic testing, we chose to err on the side of caution when interpreting direct RT-LAMP results.  
300 Furthermore, the testing landscape changed dramatically during the months we performed RT-LAMP testing. The  
301 first non-instrumented antigen test, the Abbott BinaxNOW COVID-19 Ag CARD, received FDA EUA approval in  
302 the United States on August 26, 2020 <sup>51</sup>. While the sensitivity of RT-LAMP is broadly comparable to the Abbott  
303 BinaxNOW antigen test (reported as  $1.6 \times 10^4$  -  $4.3 \times 10^4$  vRNA copies; Ct 30.3-28.8), because the former is  
304 technically straightforward and can be used as a SARS-CoV-2 diagnostic at testing sites operating under a Clinical  
305 Laboratory Improvement Amendments (CLIA) waiver, it is likely a better choice for rapid turnaround, on-site  
306 testing in most circumstances <sup>52</sup>. However, even with the existence of antigen tests, RT-LAMP surveillance  
307 programs still have a place as part of a comprehensive SARS-CoV-2 risk mitigation strategy, especially in areas  
308 where access to antigen tests is limited.

309

310 There are advantages to continuing saliva-based RT-LAMP surveillance testing. Importantly, the supply of  
311 diagnostic antigen tests remains tightly constrained, and in the United States, these tests are available only through  
312 government contracts. Widespread testing of individuals without symptoms with such a scarce resource may not be  
313 a wise use of these limited tests. Furthermore, recent studies have shown that antigen test performance may differ  
314 between asymptomatic and symptomatic populations. Compared to qRT-PCR, the sensitivity of FDA-approved  
315 antigen tests, BinaxNOW and the Quidel Sofia SARS Antigen Fluorescent Immunoassay, were 35% and 41% in  
316 asymptomatic individuals and 64% and 80% in symptomatic individuals, respectively <sup>53,54</sup>. BinaxNOW is currently  
317 only approved for use in symptomatic individuals, within 7 days of symptom onset, and samples are required to be  
318 tested within an hour of collection <sup>55</sup>. In contrast, RT-LAMP reagents do not require a government contract and can  
319 be acquired readily from commercial and non-commercial sources, and they can also be used more flexibly for  
320 surveillance purposes because RT-LAMP is not limited to use in symptomatic individuals <sup>56</sup>. Additionally, user  
321 acceptance of testing may also favor saliva-based RT-LAMP as it is less invasive than nasal swab-based tests. While  
322 an individual BinaxNOW test is rapid, performing several tests during a single day could cumulatively take as long  
323 as processing a batch of tests by RT-LAMP. For these reasons, RT-LAMP may still be the preferred testing method

324 to incorporate into a local program. In Madison, WI, two local schools have implemented RT-LAMP surveillance  
325 programs modeled on the program described here. Implementation of each program required approximately 50  
326 hours of hands-on training by our group. School staff were trained in adherence to regulations pertaining to non-  
327 diagnostic testing and to competently perform RT-LAMP assays. Each school also needed time and resources to  
328 acquire the modest lab infrastructure necessary to perform RT-LAMP. In addition, a larger saliva-based RT-LAMP  
329 surveillance program has been successfully implemented in school districts in the greater Chicago suburbs <sup>57, 58</sup>.

330

331 A looming question for both RT-LAMP and antigen testing programs is whether the real-world effectiveness of  
332 frequently testing individuals without symptoms mirrors the theoretical benefits. Several important considerations  
333 that we factored into the design of RT-LAMP testing remain true: nonsymptomatic individuals account for up to  
334 59% of all transmission (24% asymptomatic and 35% presymptomatic); low-sensitivity tests are able to effectively  
335 identify those with high levels of virus shedding, and individuals with high viral loads are likely to be responsible  
336 for a significant fraction of onward community transmission; and the duration of peak infectiousness is short, so  
337 lengthy lags in reporting test results could miss a critical window of high transmissibility <sup>10, 59</sup>. Conversely, high-  
338 quality, exceptionally well-resourced testing programs such as those at the White House and among intercollegiate  
339 athletic programs have failed to stop outbreaks <sup>60</sup>. The latter deserves special note: outbreaks in these programs  
340 occurred in spite of 100% adherence to daily testing. Data from daily sampling of individuals with incident SARS-  
341 CoV-2 infection suggests that the mean duration of time from infection to peak viral shedding is approximately  
342 three days, but some individuals potentially reach peak viral shedding in under one day <sup>61</sup>. The potential for an  
343 extremely rapid increase in viral load, which likely parallels shedding of infectious virus, means that in some cases,  
344 even daily testing might be insufficient to protect a community from someone who is newly infected.

345

346 Perhaps more importantly, the benefit of frequent testing of individuals without symptoms with RT-LAMP or other  
347 assays may be substantially undermined by risk disinhibition. When people are tested frequently, they may both  
348 underestimate their own risk of becoming infected in the interval between tests and overestimate the possibility that  
349 their similarly tested contacts are uninfected; anecdotal evidence of this phenomenon is perhaps most vividly seen  
350 in the September 26, 2020 White House Rose Garden reception for Justice Amy Coney Barrett, in which many

351 attendees were photographed not wearing masks nor following guidelines for physical distancing <sup>62</sup>. If infections  
352 among people without symptoms are rare (~0.4% of tests in this study, when combining RT-LAMP and pooled  
353 qRT-PCR positives), but 10% of the tested population views testing as license for increased risk-taking, is frequent  
354 testing of symptomless people a net positive? Appropriate messaging to the community is essential with any testing  
355 program to ensure the population understands the meaning of a test result. Such issues will require an optimization  
356 of messaging to mitigate the impact of risk disinhibition to the extent possible.

357

358 Ultimately, this study provides proof of concept and guidance for how decentralized rapid testing could be  
359 implemented in a mobile testing scenario, which may be especially useful in resource-limited settings. Despite the  
360 caveats presented above, we identified 12 SARS-CoV-2-positive individuals and likely prevented onward  
361 transmission from those individuals who otherwise would not have known they were positive. Rapid tests, although  
362 less sensitive than qRT-PCR, have shorter turnaround times and could bridge the gap between SARS-CoV-2  
363 surveillance and diagnostic testing. POC testing can be effective for identifying asymptomatic individuals but must  
364 be used in conjunction with appropriate messaging and other mitigation strategies to effectively reduce SARS-CoV-  
365 2 transmission.

366

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386

### 387 **Author Contributions**

388 CMN, MDR, RWW, DMD, CGS, DHO, SLO contributed to assay development and optimization. DMD, MTM,  
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390 EKN contributed to point of care testing and PCR confirmation. CMN, MDR, DMD, DHO contributed to data  
391 analysis, interpretation, and writing. JAK, DHO, SLO, HEB, TCF, MTM, AKH, LAH, CMC, KLH, CBB, KNF  
392 contributed to logistics and organization of point of care testing. CBB, KLH contributed to obtaining IRB and  
393 worked closely with the institutional biosafety committee on other regulatory responsibilities. MAA, ASH, WMR  
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396

### 397 **Regulatory oversight**

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636 **Figure legends**

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638 **Figure 1: Point-of-care RT-LAMP SARS-CoV-2 testing workflow. Steps 1-5.** Saliva sample preparation. **Steps**  
639 **6-7.** RT-LAMP reagent preparation. **Steps 8-10.** RT-LAMP reactions and results interpretation. A reaction color  
640 change from pink/orange to yellow after 30 minutes in at least one of two sample replicates was scored as positive.  
641 Figure was created using BioRender.com.

642

643 **Figure 2: Detection of SARS-CoV-2 in contrived saliva samples by direct RT-LAMP.** A. Representative  
644 example of a sample positive in two of two replicates. Sample is negative saliva spiked with irSARS-CoV-2. B.  
645 Representative example of a sample positive in one of two replicates C. Representative negative sample showing  
646 no colorimetric change in either replicate. D. Bar graph of results of limit of detection (LOD) assessment with  
647 contrived saliva samples from 19 volunteers. Gamma-irradiated SARS-CoV-2 (irSARS-CoV-2) vRNA load is  
648 shown as copies/ $\mu$ l on the x-axis, number of samples positive in two (black), one (dark gray), or zero (light gray)  
649 replicates is shown on the y-axis.

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651 **Figure 3: Detection of SARS-CoV-2 in 38 clinical saliva specimens by direct RT-LAMP.** The vRNA load of  
652 each clinical sample is plotted on the x-axis relative to the in-house CDC N1 qRT-PCR assay cycle threshold (Ct)  
653 on the y-axis. Black, dark gray, and light gray indicate two, one, and zero positive replicates respectively.

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663 Table 1. RT-LAMP N-gene primers

<b>Primer</b>	<b>Sequence 5'→3'</b>	<b>Concentration</b>
Outer forward primer (F3)	AACACAAGCTTTCGGCAG	0.2uM
Outer backward primer (B3)	GAAATTTGGATCTTTGTCATCC	0.2uM
Forward inner primer (FIP)	TGCGGCCAATGTTTGAATCAGCCAAGGAAATTTGGGGAC	1.6uM
Backward inner primer (BIP)	CGCATTGGCATGGAAGTCACTTTGATGGCACCTGTGTAG	1.6uM
Loop forward primer (LF)	TTCCTTGCTGATTAGTTC	0.8uM
Loop backward primer (LB)	ACCTTCGGGAACGTGGTT	0.8uM

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Table 2. RT-LAMP evaluation of SARS-CoV-2 positive clinical saliva samples.

Sample	Ct (N1 assay)	Positive by RT-LAMP	vRNA load (copies/ $\mu$ l)	Sample	Ct (N1 assay)	Positive by RT-LAMP	vRNA load (copies/ $\mu$ l)
UWHC1	27.65	0/2	3.25x10 <sup>2</sup>	UWHC20	25.80	2/2	9.48x10 <sup>2</sup>
UWHC2	32.7	0/2	10.9	UWHC21	20.18	2/2	4.40x10 <sup>4</sup>
UWHC3	20.98	2/2	5.17x10 <sup>4</sup>	UWHC22	28.92	0/2	1.13x10 <sup>2</sup>
UWHC4	24.07	2/2	3.57x10 <sup>3</sup>	UWHC23	21.26	2/2	2.10x10 <sup>4</sup>
UWHC5	26.53	2/2	6.81x10 <sup>2</sup>	UWHC24	29.92	0/2	57.2
UWHC6	30.85	1/2	37.4	UWHC25	36.71	0/2	0.796*
UWHC7	36.96	0/2	0.701	UWHC26	25.96	2/2	1.31x10 <sup>2</sup>
UWHC8	26.28	1/2	8.10x10 <sup>2</sup>	UWHC27	29.99	0/2	54.1
UWHC9	37.59	0/2	0.402	UWHC28	24.34	2/2	2.58x10 <sup>3</sup>
UWHC10	24.01	2/2	3.72x10 <sup>3</sup>	UWHC29	20.55	2/2	4.72x10 <sup>4</sup>
UWHC11	22.39	2/2	1.10x10 <sup>4</sup>	UWHC30	33.18	0/2	7.89
UWHC12	35.46	0/2	1.75	UWHC31	22.87	2/2	9.57x10 <sup>3</sup>
UWHC13	36.09	0/2	1.14	UWHC32	23.07	2/2	8.33x10 <sup>3</sup>
UWHC14	23.11	2/2	5.96x10 <sup>3</sup>	UWHC33	26.85	2/2	6.20x10 <sup>2</sup>
UWHC15	23.38	2/2	4.95x10 <sup>3</sup>	UWHC34	20.33	0/2	5.49x10 <sup>4</sup>
UWHC16	33.86	0/2	3.99	UWHC35	23	2/2	8.88x10 <sup>3</sup>
UWHC17	n/a	0/2	0	UWHC36	32.26	0/2	14.9*
UWHC18	23.02	2/2	6.34x10 <sup>3</sup>	UWHC37	33.94	0/2	4.33
UWHC19	37.31	0/2	0.612	UWHC38	25.96	2/2	1.74x10 <sup>3</sup>

\*Sample only positive in one qRT-PCR replicate.

Table 3. RT-LAMP results for 10- and 100-fold dilutions of 13 SARS-CoV-2-positive samples from UWHC.

Sample	1:10 dilution result	1:100 dilution result	Undiluted vRNA load (copies/ $\mu$ l)
UWHC1	1/2	0/2	$3.25 \times 10^2$
UWHC2	0/2	0/2	10.9
UWHC3	2/2	2/2	$5.17 \times 10^4$
UWHC4	2/2	2/2	$3.57 \times 10^3$
UWHC5	1/2	0/2	$6.81 \times 10^2$
UWHC6	0/2	0/2	37.4
UWHC7	0/2	0/2	0.701
UWHC8	1/2	0/2	$8.10 \times 10^2$
UWHC9	0/2	0/2	0.402
UWHC10	2/2	0/2	$3.72 \times 10^3$
UWHC11	2/2	1/2	$1.10 \times 10^4$
UWHC12	0/2	0/2	1.75
UWHC13	0/2	0/2	1.14

Table 4. Samples identified as potentially positive for SARS-CoV-2 by RT-LAMP during point-of-need testing.

RT-LAMP-positive sample	qRT-PCR viral load copies/ $\mu$ l
POC1	8.53
POC2	$2.15 \times 10^4$
POC3	neg
POC4	neg
POC5	neg
POC6	neg
POC7	$3.62 \times 10^5$
POC8	neg
POC9	n/a*
POC10	$2.12 \times 10^3$
POC11	neg
POC12	$1.04 \times 10^3$
POC13	$2.06 \times 10^2$
POC14	neg
POC15	52.8
POC16	$6.02 \times 10^2$
POC17	87.3
POC18	$1.17 \times 10^3$
POC19	neg
POC20	$1.38 \times 10^2$
POC21	$4.07 \times 10^2$

\*Volunteer did not consent to follow-up testing.