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 part, by testing limitations. We evaluated a simple, outdoor, mobile, colorimetric reverse 30 31 transcription loop-mediated isothermal amplification (RT-LAMP) assay workflow where self-collected 32 saliva is tested for SARS-CoV-2 RNA. From July 16 to November 19, 2020, 4,704 surveillance samples were collected from volunteers and tested for SARS-CoV-2 at 5 sites. A total of 21 samples tested 33 positive for SARS-CoV-2 by RT-LAMP; 12 were confirmed positive by subsequent quantitative reverse-34 35 transcription polymerase chain reaction (gRT-PCR) testing, while 8 were negative for SARS-CoV-2 36 RNA, and 1 could not be confirmed because the donor did not consent to further molecular testing. We 37 estimated the RT-LAMP assay's false-negative rate from July 16 to September 17, 2020 by pooling 38 residual heat-inactivated saliva that was unambiguously negative by RT-LAMP into groups of 6 or less 39 and testing for SARS-CoV-2 RNA by qRT-PCR. We observed a 98.8% concordance between the RT-40 LAMP and gRT-PCR assays, with only 5 of 421 RT-LAMP negative pools (2,493 samples) testing positive in the more sensitive gRT-PCR assay. Overall, we demonstrate a rapid testing method that can 41 be implemented outside the traditional laboratory setting by individuals with basic molecular biology 42 43 skills and can effectively identify asymptomatic individuals who would not typically meet the criteria for 44 symptom-based testing 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 performed in the United States as of February 22, 2021, yet it is estimated that 80-95% 57 of infected individuals are not tested ^{1, 2}. The availability of diagnostic testing for population surveillance 58 59 around the United States has been limited because of testing supply shortages and guidelines set by public health officials ^{3, 4}. Multiple studies have shown that asymptomatic and presymptomatic 60 individuals infected with SARS-CoV-2 can be as infectious as symptomatic individuals ^{5–9}, with recent 61 estimates of up to 59% of transmission coming from asymptomatic or presymptomatic individuals ¹⁰. 62 63 Virological assessments of SARS-CoV-2-positive individuals and coronavirus disease 2019 (COVID-64 19) patients further support the reports of asymptomatic transmission, identifying no significant differences in viral loads found in the upper respiratory tracts of asymptomatic and symptomatic 65 individuals ^{5, 7, 11–13}. Furthermore, Arons et al. (2020) demonstrated that positive viral cultures can be 66 isolated from presymptomatic patients up to 6 days before the onset of symptoms ⁵. 67

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Delays in reporting test results can prevent timely isolation of infected individuals. Most current testing 69 70 programs fail to identify and efficiently notify infected individuals. Since transmission can occur before 71 symptoms manifest, reporting delays create a major barrier to safely returning to workplaces and schools¹⁴. Therefore, there remains an urgent need for rapid tests that identify presymptomatic and 72 73 asymptomatic individuals while conserving diagnostic testing reagents. Non-diagnostic point-of-care 74 (POC) testing, used in conjunction with the current clinical diagnostic testing regimen, may improve our 75 ability to identify infectious individuals and limit their exposure to others while they are most contagious 76 and conserve clinical diagnostic tests for those who require confirmatory testing. Incorporating active 77 surveillance using POC tests as part of mitigation strategies for reopening K-12 schools could play an 78 integral role in reducing SARS-CoV-2 transmission among students, teachers and staff members, families, and the surrounding community ^{15, 16}. 79

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81 Loop-mediated isothermal amplification (LAMP) is a low-cost method for rapid target-specific detection

82 of nucleic acids ¹⁷. LAMP has long been used as an alternative to gold-standard quantitative reverse 83 transcription polymerase chain reaction (qRT-PCR) to surveil populations for a variety of pathogens, especially in resource-limited settings ¹⁸⁻²². Reverse transcription LAMP (RT-LAMP) assays have 84 recently been developed for rapid SARS-CoV-2 testing ²³⁻²⁹. RT-LAMP is an appealing candidate for 85 86 POC SARS-CoV-2 testing because it is inexpensive, circumvents supply shortages by relying on 87 different reagents than current diagnostic tests, requires minimal sample processing, and can be deployed outside of traditional laboratory settings. Recently, a number of studies have shown the 88 correlation between the presence of virus in saliva and nasopharvngeal swabs, demonstrating that 89 90 saliva specimens are a valid and reliable alternative to nasopharyngeal swab specimens for SARS-CoV-2 testing ^{30–35}. Saliva specimen self-collection is noninvasive, can be done at home, does not 91 92 require swabs or personal protective equipment, and limits direct contact between test operators and 93 testing populations. Here we describe our experience implementing a simple, rapid-turnaround, mobile, 94 non-diagnostic SARS-CoV-2 testing workflow combining self-collected saliva and RT-LAMP in 95 volunteers without symptoms of SARS-CoV-2 infection. Individuals were strongly encouraged to isolate and obtain follow-up diagnostic testing after receiving a positive result by RT-LAMP. This addresses a 96 97 key knowledge gap of how on-site RT-LAMP testing performs in real-world conditions, since virtually all 98 previous studies have only evaluated SARS-CoV-2 RT-LAMP in well-equipped molecular biology 99 laboratories.

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101 Materials and Methods

102 POC testing sites

To begin operating voluntary POC testing, we developed a system of color-coded storage bins for equipment and supplies, as well as assembled folding tables, chairs, extension cords, and coolers that could be easily decontaminated and packed to fit in a Dodge Caravan (FCA US LLC., Auburn Hills, MI) or other, similarly sized minivan for transportation between testing sites and our base laboratory facility. On July 16, 2020, we launched our first mobile POC testing sites which ultimately expanded over 18 weeks to include two workplaces, two K-12 schools, and an athletics program (Suppl. Table 1). With

109 the exception of the athletics program, sites were initially outdoors, sometimes under an overhang, but 110 otherwise open to the environment. The athletics site was a climate-controlled, indoor practice field. At 111 all sites, equipment and reagents were transported by minivan and surfaces were disinfected during 112 assembly, breakdown, and frequently throughout testing. Participant consenting and volunteer sample 113 collection were performed on-site but separated from the sample preparation and assay areas (most 114 commonly on the other side of the building). In an effort to limit contamination, each assay area was set 115 up with three separate folding tables: (1) sample heat-inactivation and preparation, (2) preparation of 116 RT-LAMP reagents and assay set-up, and (3) RT-LAMP incubation and imaging. Individuals 117 responsible for sample inactivation and performing assays wore appropriate personal protective 118 equipment (PPE) including N95 face masks, face shields or safety glasses, disposable lab coats, and 119 double gloves. In anticipation of wet and cold fall weather, by September 2020, assay workspaces were 120 transitioned to biosafety hoods in a vacant indoor laboratory space for several POC testing locations. In 121 October 2020, we received IRB approval for obtaining consent for repeat SARS-CoV-2 testing. This 122 allowed us to transition away from consenting participants at each testing time point and instead 123 allowed each enrolled participant to consent once regardless of the number of times they supplied a 124 sample. Following reports that SARS-CoV-2 RNA is stable in saliva at room temperature for prolonged 125 periods ³⁶, we also transitioned away from in-person sample collection at some of the testing sites and 126 instead distributed self-collection take-home kits for drop off at designated locations for same day 127 processing.

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129 Sample collection and preparation

We obtained approval from the University of Wisconsin-Madison Institutional Review Board (#2020-0855 and #2020-1142). Participants were advised to avoid eating, or drinking anything except for water, for 30 minutes prior to providing a sample. After providing informed consent, volunteers self-collected at least 50 µl of saliva in a 1.5 ml "safe-lock" microcentrifuge tube using a 1000 µl unfiltered pipette tip to funnel the specimen into the tube. Each volunteer disinfected the outside of the tube with a premoistened disinfectant wipe. Samples collected in-person were typically processed within 3 hours of

136 collection through our RT-LAMP mobile testing workflow, while samples collected using take-home kits 137 were typically processed within 30 hours (Figure 1). Samples were first incubated in a heat block at 138 65°C for 30 minutes to inactivate SARS-CoV-2 ³⁷ and then incubated in another preset heat block at 139 98°C for 3 minutes to improve nucleic acid detection and inactivate salivary enzymes ³⁸. The inactivated 140 saliva was then centrifuged for 2 minutes in a benchtop microcentrifuge. Fifty microliters of the saliva 141 supernatant were then added to 50 µl of 1x phosphate buffered saline, pH 7.4 (1x PBS).

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143 RT-LAMP reactions

144 Three microliters of the saliva/PBS mixture for each sample was added in duplicate to 17 µl of a 145 colorimetric RT-LAMP reaction mix containing WarmStart colorimetric LAMP mastermix (NEB, 146 catalogue# M1800), water, and a set of six SARS-CoV-2-specific RT-LAMP primers designed against the N gene ³⁸. The SARS-CoV-2 RT-LAMP primer set was previously designed by Broughton et al. and 147 148 is currently used in an FDA emergency use authorized (EUA) COVID-19 test by Color Genomics (Table 1) ^{39, 40}. Reactions were incubated for 30 minutes at 65°C. A smartphone or tablet was used to record 149 150 images of each reaction before (time = 0) and after the incubation period (time = 30). A color change 151 from pink/orange to yellow in at least 1 of 2 replicates was scored relative to gamma-irradiated SARS-152 CoV-2 (irSARS-CoV-2, BEI Resources, Manassas, VA) that was directly added to RT-LAMP reactions 153 as a positive control in each batch of reactions at concentrations ranging from 220-3,333 copies/ul (2.2x10⁵ - 3.33x10⁶ copies/ml). irSARS-CoV-2 was diluted and aliquoted as ready-to-run positive 154 155 control standards and stored at -80°C. On the day of testing, the positive controls were removed from 156 the freezer and stored on ice at POC sites. Individuals whose samples were recorded as potentially positive for SARS-CoV-2 by RT-LAMP were contacted by an infectious disease clinician in accordance 157 158 with the IRB protocol and urged to obtain a clinical diagnostic test to confirm findings and self-isolate in 159 accordance with public health recommendations.

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

162 To estimate the limit of detection of the RT-LAMP assay, contrived positive saliva samples were

163 prepared by adding irSARS-CoV-2 diluted from 1x10⁴-10 copies/µl (1x10⁷-1x10⁴copies/ml) or from $5x10^{4}$ -50 copies/µl ($5x10^{7}$ - $5x10^{4}$ copies/ml) directly into unaltered saliva collected from a total of 25 164 165 SARS-CoV-2-negative individuals. Dilutions were based on two independent, in-house gRT-PCR experiments showing that the ir-SARS-CoV-2 stock concentration ranged from 7.89x10⁶ - 8.23x10⁶ 166 copies/µl (7.89x10⁹ - 8.23x10⁹ copies/ml). In two RT-LAMP experiments, four serial dilutions of irSARS-167 CoV-2 were prepared for each saliva sample in duplicate. RT-LAMP reactions were set up as described 168 169 previously. Negative controls consisting of 1x PBS and positive controls consisting of 1x10⁴ copies/µl (1x10⁷ copies/ml) irSARS-CoV-2 in water were also prepared in duplicate. Reactions were called 170 171 positive if a color change from pre-amplification to post-amplification occurred in at least 1 of 2 172 replicates that was consistent with that of the positive controls.

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174 Limit of detection (LOD) estimation using clinical samples

175 De-identified discard saliva samples from 38 SARS-CoV-2-positive patients were provided by the 176 University of Wisconsin Hospitals and Clinics (UWHC) for evaluation of RT-LAMP performance with 177 known positive saliva samples. Clinical saliva samples were originally collected and stored at 4°C for up 178 to 4 weeks prior to assessment by RT-LAMP. Additional 10-fold and 100-fold dilutions were prepared 179 for 13 of the samples in saliva collected from a negative volunteer. Clinical samples and dilutions were 180 prepared as described previously except that 20-50 µl of heat-inactivated sample, dependent on total 181 sample volume, was added to an equal volume of 1x PBS in a clean 1.5 ml screw-top tube and pipetted 182 gently to mix. For each sample, 3 µl was then added to duplicate colorimetric RT-LAMP reactions. 183 Negative and positive control reactions (described previously) were also prepared in duplicate except 184 that saliva collected from a negative volunteer was used as the negative control for these reactions. 185 RT-LAMP reactions were prepared and images collected as described previously.

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187 Quantitative RT-PCR

188 POC samples

189 We measured vRNA concentration using sensitive qRT-PCR in a subset of the inactivated saliva 190 samples described above after initial evaluation using RT-LAMP. Saliva samples that were negative for 191 SARS-CoV-2 by RT-LAMP were pooled into groups of 6 or fewer for gRT-PCR to balance cost 192 effectiveness with reasonable estimated detection sensitivity. Ten additional, individual RT-LAMP-193 negative samples were submitted as negative controls alongside samples identified as positive by RT-194 LAMP. Saliva samples that were identified as positive for SARS-CoV-2 by RT-LAMP were tested by 195 gRT-PCR individually to estimate our POC LOD. RNA was isolated from up to 150 µl saliva and 196 combined with an equivalent volume of nuclease-free water using the Viral Total Nucleic Acid kit for the 197 Maxwell RSC instrument (Promega, Madison, WI) following the manufacturer's instructions. Viral load 198 quantification was performed using a sensitive gRT-PCR assay developed by the CDC to detect SARS-199 CoV-2 (specifically the N1 assay) and commercially available from IDT (Coralville, IA). The assay was 200 run on a LightCycler 96 or LC480 instrument (Roche, Indianapolis, IN) using the Tagman Fast Virus 1-201 step Master Mix enzyme (Thermo Fisher, Waltham, MA). The limit of detection of this assay is 202 estimated to be 0.2 genome equivalents/µl (200 genome equivalents/ml) saliva. To determine the vRNA load, samples were interpolated onto a standard curve consisting of serial 10-fold dilutions of in 203 204 vitro transcribed SARS-CoV-2 N gene RNA kindly provided by Nathan Grubaugh (Yale University) and 205 described by Dudley et al. ³⁵.

206

207 Clinical samples

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

- 214
- 215 Results

216 LOD estimation using contrived saliva samples

We assessed the LOD for minimally processed saliva samples collected from 25 volunteers over two 217 RT-LAMP experiments using irSARS-CoV-2 spiked into negative saliva samples (Figure 2A and 2B). In 218 our first experiment (S1-S3), we detected irSARS-CoV-2 in at least 1 of 2 replicates at 1x10² copies/µl 219 (1x10⁵ copies/ml) in all 3 samples (Figure 2A). In our second experiment (S4-S25), we detected 220 221 irSARS-CoV-2 by RT-LAMP in 2/2 replicates at 5x10⁴ copies/µl (5x10⁷ copies/ml) for 95% of samples, at 5x10³ copies/µl (5x10⁶ copies/ml) for 62% of samples, and at 500 copies/µl (5x10⁵ copies/ml) for 10% 222 223 of samples. When we included samples called positive in at least 1 of 2 replicates (see Methods), the 224 percentage of contrived samples positive by RT-LAMP at each of the aforementioned dilutions were 100%, 90%, and 33.3% respectively (Figure 2B). One sample was omitted from the analysis because it 225 turned yellow before the RT-LAMP reaction incubation began and was therefore uninterpretable. 226 227 Because in POC testing we defined a positive RT-LAMP result as an observed post-incubation color change to yellow in at least 1 replicate, these results suggested that our 90% LOD is between 1x10² 228 and $5x10^3$ copies/µl ($1x10^5 - 5x10^6$ copies/ml). 229

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231 LOD estimation using clinical samples

To assess the performance of SARS-CoV-2 RT-LAMP in known SARS-CoV-2 positive saliva samples 232 233 as opposed to contrived positive samples, we acquired deidentified, discarded saliva samples collected 234 from 38 patients with laboratory confirmed SARS-CoV-2 from UWHC. Nineteen of 38 undiluted saliva samples were positive for SARS-CoV-2 in 2/2 replicates by RT-LAMP (Figure 3; Table 2). Two 235 236 additional samples were positive in 1 of 2 replicates. Quantitative RT-PCR data showed that the viral RNA (vRNA) loads of the positive samples ranged from 131 copies/µl to 5.7x10⁴ copies/µl (1.31x10⁵-237 5.71x10⁷ copies/ml) which was consistent with our LOD range for contrived samples (Table 3). 238 239 Furthermore, for the 13 samples diluted 10-fold and 100-fold, detection decreased with increasing 240 dilution factor (Table 4).

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242 POC SARS-CoV-2 RT-LAMP testing

243 From July 16 to November 19, 2020, SARS-CoV-2 RT-LAMP was used to test a total of 4,704 samples 244 collected from 5 locations. Participants were enrolled into the study regardless of their SARS-CoV-2 245 symptom status on the day of testing. Seventy-one percent of the samples were obtained from 246 individuals at two research facilities, 11% from two K-12 schools, and 18% from an athletics program 247 (Supplemental Table 1). A total of 21 samples were identified as positive for SARS-CoV-2 by RT-LAMP 248 based on a colorimetric change from pink/orange to yellow in at least 1 of 2 sample replicates. Similar 249 to our experience with our contrived LOD samples, about 0.40% (19/4,704) of samples collected during 250 POC testing exhibited a color change to vellow prior to RT-LAMP assay amplification and were 251 therefore uninterpretable. Follow up qRT-PCR testing was conducted on each sample that appeared 252 positive after the 30 minute amplification reaction throughout the study to determine vRNA load. Twelve of the 21 samples called positive in RT-LAMP had detectable SARS-CoV-2 RNA by gRT-PCR. Viral 253 RNA loads of these samples ranged from 8.58 copies/µl to 3.62x10⁵ copies/µl (8.58x10³ copies/ml-254 3.62×10^8 copies/ml) with a median of 504.5 copies/µl (5.04×10^5 copies/ml) (Table 4). Eight of the saliva 255 256 samples identified as positive by RT-LAMP were negative by gRT-PCR, suggesting that they were 257 false-positive RT-LAMP results. One RT-LAMP-positive sample was not tested by gRT-PCR because 258 the participant did not consent to additional molecular testing. For volunteers who consented to 259 additional research testing from July 16 to September 17, gRT-PCR testing was conducted for pools of 260 6 or fewer for all residual, heat-inactivated samples that appeared unambiguously negative by RT-261 LAMP. A total of 421 RT-LAMP-negative pools (2,493 samples) were tested to estimate the number of 262 SARS-CoV-2-positive samples missed by RT-LAMP. Quantitative RT-PCR detected SARS-CoV-2 263 nucleic acids in 5 pools of RT-LAMP-negative samples. Four out of five of the positive pools contained 264 levels of SARS-CoV-2 that were below the estimated LOD range for RT-LAMP using crude samples with vRNA load estimates of 0.236, 0.444, 0.460, 37.5, and 142 copies/ul (236, 444, 460, 3.75x10⁴, and 265 1.42x10⁵ copies/ml). Taken together, the low prevalence of SARS-CoV-2 in our volunteer testing 266 267 population (0.36%, including RT-LAMP-negative, gRT-PCR-positive pools) and the low vRNA load of 268 pools positive by follow-up qRT-PCR, suggest that these 5 pools likely contained only a single positive 269 sample each and suggests a false-negative rate of 0.02% (5/2.493) (Table 4).

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271 Discussion

272 Strategic surveillance testing of asymptomatic individuals has been suggested as an important 273 mitigation strategy for places at high risk for close contact, indoor SARS-CoV-2 transmission: schools, 274 workplaces, places of worship, and prisons, among others. Decentralized, mobile RT-LAMP-based 275 POC testing workflows can provide same-day results which can enable people with potential SARS-276 CoV-2 infections to quickly self-isolate and then obtain confirmatory diagnostic testing. The low per-test 277 cost (approximately \$7 per sample tested in duplicate) allows for repeated testing to identify incident 278 infections and reduce the duration of a potentially infected individual's exposure to others. While RT-279 LAMP is not as sensitive as diagnostic gRT-PCR tests in laboratory testing, gRT-PCR tests require 280 centralized labs, which in turn leads to lengthy turnaround times. Over a period of 18 weeks, we 281 performed 4,704 SARS-CoV-2 tests across 5 sites using a simple, saliva-based, direct RT-LAMP 282 assay. This work demonstrates the scalability of decentralized, mobile RT-LAMP-based testing and 283 addresses a key knowledge gap of how POC RT-LAMP testing performs outside of well-equipped 284 molecular biology laboratories.

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286 Our initial experiments using direct RT-LAMP with contrived saliva samples from a total of 25 donors demonstrated a LOD that ranged from 1x10² copies/µl (100% in at least 1 replicate for S1-S3) to 5x10³ 287 288 copies/µl (90% in at least one replicate for S4-S25). Taken together, these data suggest that the actual 289 LOD for RT-LAMP without RNA isolation may be dependent on the individual sample due to heterogeneity of saliva pH and composition ⁴¹⁻⁴³. Overall, the RT-LAMP results for 38 clinical saliva 290 291 samples obtained from SARS-CoV-2-positive individuals at the UWHC, were consistent with those for 292 the contrived samples. We recognize that more clinical samples are required for a comprehensive 293 clinical validation, but the LOD observed in clinical samples is further supported by the low vRNA loads 294 obtained from gRT-PCR-confirmed SARS-CoV-2-positive samples identified in our volunteer population 295 (Table 4). The performance of our RT-LAMP POC testing workflow demonstrates that inexpensive, 296 mobile testing can be successfully performed outdoors or in other non-traditional laboratory settings to

297 identify SARS-CoV-2-positive individuals regardless of whether or not symptoms are present. Our 298 observed SARS-CoV-2 RT-LAMP positivity rate was 0.25% (12/4,704) for samples confirmed by follow-299 up gRT-PCR. Interestingly, the positivity rate of 0.25% in our volunteer population was lower than 300 expected given the disease activity in our region during this period of time was listed as "critically high", 301 particularly between September 1 and November 19, 2020 when the county had a 5.42% positivity rate (19,031 positive tests out of 350,722)^{44, 45}. The low positivity rate in our volunteer population may be 302 303 partly explained by the fact that 71% of tested saliva specimens came from two research facilities 304 where mask wearing and physical distancing guidelines were implemented early in the pandemic and 305 followed relatively stringently (Supplemental Table 1). Volunteers for nonsymptomatic research testing 306 might also have a different risk profile from the overall population.

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308 Potential drawbacks of colorimetric RT-LAMP-based surveillance for SARS-CoV-2 as described here 309 include the fact that minimally-processed saliva can result in variable reaction color change without the 310 presence of the target RNA. However, modifications of RT-LAMP-based SARS-CoV-2 assays to 311 reduce saliva sample variability, improve result ambiguity, and increase throughput have recently been 312 reported elsewhere and may improve the implementation of RT-LAMP-based assays for POC use ^{46–50}. 313 In addition, we relied on a manual RT-LAMP format during POC testing. Reading assays "by eye" 314 inevitably results in a somewhat subjective determination of positives. Reducing false-positive results in 315 our POC volunteer populations required consistent use of duplicate reactions for each individual, which 316 reduced assay throughput and increased the per-sample cost. Furthermore, the testing landscape 317 changed dramatically during the months we performed RT-LAMP testing. The first non-instrumented antigen test, the Abbott BinaxNOW COVID-19 Ag CARD, received FDA EUA approval in the United 318 States on August 26, 2020⁵¹. While the sensitivity of RT-LAMP is broadly comparable to the Abbott 319 BinaxNOW antigen test (reported as 1.6×10^4 - 4.3×10^4 vRNA copies; Ct 30.3-28.8), because the former 320 321 is technically straightforward and can be used as a SARS-CoV-2 diagnostic at testing sites operating 322 under a Clinical Laboratory Improvement Amendments (CLIA) waiver, it is likely a better choice for rapid turnaround, on-site testing in most circumstances ⁵². However, even with the existence of antigen 323

tests, RT-LAMP surveillance programs still have a place as part of a comprehensive SARS-CoV-2 risk
 mitigation strategy, especially in areas where access to antigen tests is limited.

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327 There are advantages to continuing saliva-based RT-LAMP surveillance testing. Importantly, the supply 328 of diagnostic antigen tests remains tightly constrained, and in the United States, these tests are 329 available only through government contracts. Widespread testing of individuals without symptoms with 330 such a scarce resource may not be a wise use of these limited tests. Furthermore, recent studies have 331 shown that antigen test performance may differ between asymptomatic and symptomatic populations. 332 Compared to qRT-PCR, the sensitivity of FDA-approved antigen tests, BinaxNOW and the Quidel Sofia 333 SARS Antigen Fluorescent Immunoassay, were 35% and 41% in asymptomatic individuals and 64% and 80% in symptomatic individuals, respectively ^{53, 54}. BinaxNOW is currently only approved for use in 334 335 symptomatic individuals, within 7 days of symptom onset, and samples are required to be tested within an hour of collection ⁵⁵. In contrast, RT-LAMP reagents do not require a government contract and can 336 337 be acquired readily from commercial and non-commercial sources, and they can also be used more flexibly for surveillance purposes because RT-LAMP is not limited to use in symptomatic individuals ⁵⁶. 338 339 Additionally, user acceptance of testing may also favor saliva-based RT-LAMP as it is less invasive 340 than nasal swab-based tests. While an individual BinaxNOW test is rapid, performing several tests 341 during a single day could cumulatively take as long as processing a batch of tests by RT-LAMP. For 342 these reasons, RT-LAMP may still be the preferred testing method to incorporate into a local program. 343 In Madison, WI, two local schools have implemented RT-LAMP surveillance programs modeled on the 344 program described here. Implementation of each program required approximately 50 hours of hands-on 345 training by our group. School staff were trained in adherence to regulations pertaining to non-diagnostic 346 testing and to competently perform RT-LAMP assays. Each school also needed time and resources to 347 acquire the modest lab infrastructure necessary to perform RT-LAMP. In addition, a larger saliva-based 348 RT-LAMP surveillance program has been successfully implemented in school districts in the greater 349 Chicago suburbs 57, 58.

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351 A looming question for both RT-LAMP and antigen testing programs is whether the real-world 352 effectiveness of frequently testing individuals without symptoms mirrors the theoretical benefits. Several 353 important considerations that we factored into the design of RT-LAMP testing remain true: 354 nonsymptomatic individuals account for up to 59% of all transmission (24% asymptomatic and 35% 355 presymptomatic): low-sensitivity tests are able to effectively identify those with high levels of virus 356 shedding, and individuals with high viral loads are likely to be responsible for a significant fraction of 357 onward community transmission; and the duration of peak infectiousness is short, so lengthy lags in reporting test results could miss a critical window of high transmissibility ^{10, 59}. Conversely, high-quality, 358 359 exceptionally well-resourced testing programs such as those at the White House and among intercollegiate athletic programs have failed to stop outbreaks ⁶⁰. The latter deserves special note: 360 361 outbreaks in these programs occurred in spite of 100% adherence to daily testing. Data from daily 362 sampling of individuals with incident SARS-CoV-2 infection suggests that the mean duration of time 363 from infection to peak viral shedding is approximately three days, but some individuals potentially reach 364 peak viral shedding in under one day ⁶¹. The potential for an extremely rapid increase in viral load, 365 which likely parallels shedding of infectious virus, means that in some cases, even daily testing might 366 be insufficient to protect a community from someone who is newly infected.

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368 Perhaps more importantly, the benefit of frequent testing of individuals without symptoms with RT-369 LAMP or other assays may be substantially undermined by risk disinhibition. When people are tested 370 frequently, they may both underestimate their own risk of becoming infected in the interval between 371 tests and overestimate the possibility that their similarly tested contacts are uninfected; anecdotal 372 evidence of this phenomenon is perhaps most vividly seen in the September 26, 2020 White House 373 Rose Garden reception for Justice Amy Coney Barrett, in which many attendees were photographed not wearing masks nor following guidelines for physical distancing ⁶². If infections among people without 374 375 symptoms are rare (~0.4% of tests in this study, when combining RT-LAMP and pooled gRT-PCR 376 positives), but 10% of the tested population views testing as license for increased risk-tasking, is 377 frequent testing of symptomless people a net positive? Appropriate messaging to the community is

essential with any testing program to ensure the population understands the meaning of a test result.
Such issues will require an optimization of messaging to mitigate the impact of risk disinhibition to the
extent possible.

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382 Ultimately, this study provides proof of concept and guidance for how decentralized rapid testing could 383 be implemented in a mobile testing scenario, which may be especially useful in resource-limited 384 settings. Despite the caveats presented above, we identified 12 SARS-CoV-2-positive individuals and 385 likely prevented onward transmission from those individuals who otherwise would not know they were 386 positive. Rapid tests, although less sensitive than qRT-PCR, have shorter turnaround times and could 387 bridge the gap between SARS-CoV-2 surveillance and diagnostic testing. POC testing can be effective 388 for identifying asymptomatic individuals but must be used in conjunction with appropriate messaging 389 and other mitigation strategies to effectively reduce SARS-CoV-2 transmission.

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413 Author Contributions

414 CMN, MDR, RWW, DMD, CGS, DHO, SLO contributed to assay development and optimization. DMD, 415 MTM, RWW, CMN, MRS, AMW, MIB, KNF, MDR, LAH, OEH, RVM, CMC, SLO, MRR, TCF, TMP, 416 EDS, LMS, EKN contributed to point of care testing and PCR confirmation. CMN, MDR, DMD, DHO 417 contributed to data analysis, interpretation, and writing. JAK, DHO, SLO, HEB, TCF, MTM, AKH, LAH, 418 CMC, KLH, CBB, KNF contributed to logistics and organization of point of care testing. CBB, KLH contributed to obtaining IRB and worked closely with the institutional biosafety committee on other 419 420 regulatory responsibilities. MAA, ASH, WMR contributed to providing residual SARS-CoV-2 positive 421 saliva samples and sample information from the University of Wisconsin Hospitals and Clinics. All 422 authors contributed to editing the manuscript.

423

424 Regulatory oversight

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681 Figure 2: Detection of SARS-CoV-2 in contrived saliva samples by direct RT-LAMP. A. Initial limit 682 of detection (LOD) assessment with contrived saliva samples from 3 volunteers (S1, S2, S3). RT-LAMP 683 reactions determined to be negative are pink and those determined to be positive are yellow. 684 Quantitative RT-PCR vRNA loads are presented as copies/µl above the replicates for each sample. B. Bar graph showing an expanded assessment of RT-LAMP LOD for 22 additional contrived saliva 685 686 samples (S4-S25). Gamma-irradiated SARS-CoV-2 (irSARS-CoV-2) vRNA load is shown as copies/µl 687 on the x-axis, number of samples positive in 2 (black), 1 (dark gray), or 0 (light gray) replicates is shown 688 on the y-axis.

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| 690 | Figure 3: Detection of SARS-CoV-2 in 38 clinical saliva specimens by direct RT-LAMP. The vRNA |
|-----|---|
| 691 | load of each clinical sample is plotted on the x-axis relative to the in-house CDC N1 qRT-PCR assay |
| 692 | cycle threshold (Ct) on the y-axis. Black, dark gray, and light gray indicate 2, 1, and 0 positive replicates |
| 693 | respectively. |
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702 Table 1. RT-LAMP N-gene primers

| Primer | Sequence 5'->3' | Concentration |
|-----------------------------|---|---------------|
| Outer forward primer (F3) | AACACAAGCTTTCGGCAG | 0.2uM |
| Outer backward primer (B3) | GAAATTTGGATCTTTGTCATCC | 0.2uM |
| Forward inner primer (FIP) | TGCGGCCAATGTTTGTAATCAGCCAAGGAAATTTTGGGGAC | 1.6uM |
| Backward inner primer (BIP) | CGCATTGGCATGGAAGTCACTTTGATGGCACCTGTGTAG | 1.6uM |
| Loop forward primer (LF) | TTCCTTGTCTGATTAGTTC | 0.8uM |
| Loop backward primer (LB) | ACCTTCGGGAACGTGGTT | 0.8uM |

medRxiv preprint doi: https://doi.org/10.1101/2020.07.28.20164038; this version posted February 28, 2021. The copyright holder for this preprint (which was not certified by peer review) is the author/funder, who has granted medRxiv a license to display the preprint in perpetuity. Table 2. RT-LAMP evaluation of SARS-2010-2 positive clinical salivariant preprint in perpetuity.

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

*Sample only positive in one qRT-PCR replicate.

| Sample | 1:10 dilution result | 1:100 dilution result | Undiluted vRNA load (copies/µl) |
|--------|----------------------|-----------------------|---------------------------------|
| UWHC1 | 1/2 | 0/2 | 3.25x10 ² |
| UWHC2 | 0/2 | 0/2 | 10.9 |
| UWHC3 | 2/2 | 2/2 | 5.17x10 ⁴ |
| UWHC4 | 2/2 | 2/2 | 3.57x10 ³ |
| UWHC5 | 1/2 | 0/2 | 6.81x10 ² |
| UWHC6 | 0/2 | 0/2 | 37.4 |
| UWHC7 | 0/2 | 0/2 | 0.701 |
| UWHC8 | 1/2 | 0/2 | 8.10x10 ² |
| UWHC9 | 0/2 | 0/2 | 0.402 |
| UWHC10 | 2/2 | 0/2 | 3.72x10 ³ |
| UWHC11 | 2/2 | 1/2 | 1.10x10 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/µl |
|-------------------------|------------------------------|
| POC1 | 8.53 |
| POC2 | 2.15x10 ⁴ |
| POC3 | neg |
| POC4 | neg |
| POC5 | neg |
| POC6 | neg |
| POC7 | 3.62x10 ⁵ |
| POC8 | neg |
| POC9 | n/a* |
| POC10 | 2.12x10 ³ |
| POC11 | neg |
| POC12 | 1.04x10 ³ |
| POC13 | 2.06x10 ² |
| POC14 | neg |
| POC15 | 52.8 |
| POC16 | 6.02x10 ² |
| POC17 | 87.3 |
| POC18 | 1.17x10 ³ |
| POC19 | neg |
| POC20 | 1.38x10 ² |
| POC21 | 4.07x10 ² |

*Volunteer did not consent to follow-up testing.

Saliva preparation



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