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1 Harnessing recombinase polymerase amplification for 2 rapid detection of SARS-CoV-2 in resource-limited 3 settings

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15 **Abstract**

16 The COVID-19 pandemic has challenged testing capacity worldwide. The mass testing
17 needed to stop the spread of the virus requires new molecular diagnostic tests that are faster
18 and with reduced equipment requirement, but as sensitive as the current gold standard
19 protocols based on polymerase chain reaction.

20 We developed a fast (25-35 minutes) molecular test using reverse transcription recombinase
21 polymerase amplification for simultaneous detection of two conserved regions of the virus,
22 targeting the E and RdRP genes. The diagnostic platform offers two complementary detection
23 methods: real-time fluorescence or visual dipstick.

24 The analytical sensitivity of the test by real-time fluorescence was 9.5 (95% CI: 7.0-18) RNA
25 copies per reaction for the E gene and 17 (95% CI: 11-93) RNA copies per reaction for the
26 RdRP gene. The analytical sensitivity for the dipstick readout was 130 (95% CI: 82-500)
27 RNA copies per reaction. The assay showed high specificity with both detection methods
28 when tested against common seasonal coronaviruses, SARS-CoV and MERS-CoV model
29 samples. The dipstick readout demonstrated potential for point-of-care testing, with simple or
30 equipment-free incubation methods and a user-friendly prototype smartphone application was
31 proposed with data capture and connectivity.

32 This ultrasensitive molecular test offers valuable advantages with a swift time-to-result and it
33 requires minimal laboratory equipment compared to current gold standard assays. These
34 features render this diagnostic platform more suitable for decentralised molecular testing.

35 **Keywords:** molecular testing, isothermal amplification, recombinase polymerase
36 amplification, SARS-CoV-2.

37 **Introduction**

38 At the end of December 2019, a public health alert was released from Wuhan, Hubei
39 province, in China reporting cases of “viral pneumonia of unknown cause” observed in
40 several patients with severe acute respiratory syndrome¹. Eventually, the newly identified
41 virus was designated as severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)² and
42 the disease caused by the virus was named COVID-19³. As of 8th January 2021, a year after
43 the discovery of the human coronavirus SARS-CoV-2, the World Health Organization
44 reported globally over 86.4 million confirmed cases and 1.8 million deaths from COVID-19⁴.
45
46 Rapid development of diagnostic tests for detection of SARS-CoV-2 has been vital to limit
47 the spread of the virus⁵. Molecular diagnosis is necessary to identify patients actively infected
48 when COVID-19 symptoms are not clearly differentiable from other coronaviruses, for
49 instance HCoV-NL63, HCoV-OC43 and HCoV-229E, causing common cold, or the deadly
50 Severe Acute Respiratory Syndrome Coronavirus (SARS-CoV) and Middle East Respiratory
51 Syndrome Coronavirus (MERS-CoV). These tests are also needed to identify asymptomatic
52 cases⁶ (not showing symptoms) or pre-symptomatic cases⁷ (not showing symptoms at the
53 time of test but developing symptoms later) which can be infectious. Since the early stages of
54 the pandemic, the World Health Organization recommended the use of quantitative reverse
55 transcription polymerase chain reaction (qRT-PCR) for nucleic acid amplification as the gold
56 standard diagnostic for SARS-CoV-2⁸. Some in-house qRT-PCR protocols were swiftly
57 developed and recommended for wide use in reference laboratories, such as Hong Kong
58 University (HKU)⁹, Charité Institute of Virology Universitätsmedizin Berlin (Charité-
59 Berlin)¹⁰ and United States CDC (US CDC)¹¹. Although targeting various conserved regions
60 of SARS-CoV-2, these qRT-PCR protocols all function with multiple targets to make the test
61 more sensitive and specific (Supplementary Fig. 1).

62

63 Despite the World Health Organisation's recommendation to use qRT-PCR technology for
64 detection of SARS-CoV-2, the pandemic has highlighted major issues in relying on only one
65 technology: a worldwide shortage of qRT-PCR reagents and instruments considerably slowed
66 down testing^{12,13}. The massive number of tests needed to contain the spread of the virus could
67 not be met for many months, even in high-income countries¹⁴. Having viable alternatives to
68 qRT-PCR for acute COVID-19 that are as sensitive, but faster and simpler to use –
69 particularly in decentralised and resource-limited settings in the low and middle income
70 countries – could increase the testing capacity and reduce community transmission¹⁵.

71

72 Recombinase polymerase amplification (RPA), loop-mediated isothermal amplification
73 (LAMP) and other isothermal amplification methods could meet the needs for mass
74 virological testing as alternatives to PCR¹⁶. They are portable, faster, giving results in less
75 than 30 minutes compared to several hours with qRT-PCR, enabling testing in settings with
76 scarce resources, without a thermocycler^{17,18}. Indeed, the first molecular test to receive FDA
77 authorisation for emergency use, which can be operated by the patient without going to a test
78 centre, harnesses isothermal amplification (RT-LAMP) to deliver an easy-to-use device¹⁹.
79 Several other RPA-based tests for SARS-CoV-2 reported in the literature present good
80 performances^{20,21}.

81

82 Reverse transcription RPA (RT-RPA) can rapidly amplify viral RNA and detection of the
83 RPA product has been demonstrated by several methods, the most common are by real-time
84 fluorescence or using dipsticks. An ultrasensitive diagnostic assay using RPA and dipstick
85 was successfully demonstrated by our group for HIV using novel nanoparticles²². While
86 LAMP has the advantage of being a patent-free method, it operates at higher temperatures

87 (60-65°C compared to 37-42°C for RPA) and it requires two pairs of primers instead of one,
88 making its design slightly more complex. Choosing an RPA-based amplification enables the
89 reaction to be carried out with basic equipment.

90

91 Mobile phone-based diagnostic tests can help reduce the need for specific equipment, whilst
92 providing mobile connectivity, which has emerged as an important criteria in REASSURED
93 diagnostic tests²³. By taking advantage of their processing speeds, display, storage capacity,
94 their high resolution camera and connectivity, smartphones are useful devices to store
95 information relative to the test and the patient, analyse test results with an enhanced readout
96 compared to the naked eye, to communicate the result to a local hospital and send alerts in
97 case a new outbreak or cluster is detected^{24,25}.

98

99 Here we report the development of a rapid molecular diagnostic for the detection of SARS-
100 CoV-2 by RT-RPA (Fig. 1a) simultaneously detecting two targets (Fig. 1b). We aimed to
101 design two alternative readouts which are both multiplexed: real-time fluorescence (Fig. 1c)
102 and dipsticks (Fig. 1d). Offering two detection methods makes the assay more accessible to
103 different settings, depending on their resources. The multiplexing of the targets has also been
104 used by gold standard qPCR methods (Supplementary Fig. 1) as it offers a greater confidence
105 on the test result, which can be confirmed if only one gene was detected.

106

107 **Fig. 1 Schematic representation of the rapid and multiplex RT-RPA assay with real-**
108 **time fluorescence and dipstick detection.**

109

110 **Methods**

111 **Reagents and equipment**

112 RPA primers and the cDNA control for the N gene were obtained from Integrated DNA
113 Technologies. The cDNA controls for the E, RdRP and Orflab genes were supplied by
114 GenScript Biotech (pUC57-2019-nCoV-PC:E, pUC57-2019-nCoV-PC:RdRP, pUC57-2019-
115 nCoV-PC:ORF1ab). The coronavirus specificity panel (SARS-CoV-2, SARS-CoV, MERS-
116 CoV, HCoV-NL63, HCoV-OC43 and HCoV-229E) was obtained from the European Virus
117 Archive (EVAg). These RNA samples were supplied as full-length virus RNA with reported
118 cycle threshold (Ct) values between 28-30 from qRT-PCR assays. The fluorescent probes
119 were synthesized by Eurogentec. The RPA reactions kits were ordered from TwistDX. The
120 QIAquick Gel Extraction and PCR Purification kits were ordered from Qiagen. The
121 Phusion™ High-Fidelity DNA Polymerase kit and the M-MLV reverse transcriptase were
122 from Thermo Fisher Scientific. The *in vitro* transcription and the RNA purification was
123 performed with the HiScribe™ T7 Quick High Yield RNA Synthesis kit and Monarch RNA
124 CleanUp kit from New England Biolabs Ltd. SUPERase•In™ RNase inhibitor from
125 Invitrogen was added to the RNA standards. Human saliva from healthy and pooled donors
126 (cat. 991-05-P-PreC) was purchased from Lee Biosolutions, Inc. cDNA concentrations were
127 measured on a Nanodrop™ One/One^C microvolume UV-Vis spectrophotometer. RNA
128 concentrations were measured on a Qubit 4 fluorometer using the Qubit™ RNA HS Assay
129 Kit (Invitrogen). Fluorescent readings were done on the microplate reader SpectraMax® iD3
130 from Molecular Devices, for initial screening of SARS-CoV-2 genes, then the Axxin® T16-
131 ISO was used for the duplex diagnostic platform. The dipsticks and running buffer were
132 obtained from Abingdon Health.

133

134 **Screening of four different genes (N, E, RdRP and Orflab genes) by real-time RPA**

135 RPA primers and probes with a FAM fluorophore were designed for four targets
136 (Supplementary Table 1) and screened using the TwistAmp® exo RPA reactions. The 50 µL

137 reactions contained TwistAmp® exo RPA pellets resuspended in 29.5 µL Rehydration Buffer
138 (TwistDX), 2.1 µL of forward primer (at concentration 10 µM), 2.1 µL of reverse primer (at
139 concentration 10 µM), 0.6 µL of probe (at concentration 10 µM), 1 µL of corresponding
140 cDNA template and 12.2 µL of nuclease-free water. Finally, 2.5 µL of magnesium acetate (at
141 concentration 280 mM) was added to start the reaction. Three cDNA concentrations (50, 500,
142 5000 copies) were tried along with a non-template control (NTC). The reactions were
143 incubated at 39°C for 30 minutes and real-time fluorescence was recorded using a microplate
144 reader (excitation wavelength 495 nm and emission 520 nm). The screen was done in
145 technical replicates (N=2). Background correction was done on all data by subtracting the
146 measurement value taken at ≈ 60 seconds from all other measurement values, and the values
147 before 60 seconds were set to zero. Then, the average values of duplicates were plotted on
148 GraphPad Prism along with error bars, corresponding to the standard deviation. The
149 fluorescence threshold value for the RPA screen of the four genes with cDNA was set to
150 25,000. This threshold value was roughly calculated by averaging fluorescence signals from
151 several NTC reactions and adding 3 times the associated standard deviation. The average
152 time to threshold, defined as the time corresponding to the intersection of the amplification
153 curve with the threshold value, was determined for each gene.

154

155 **Synthesis of RNA standards for SARS-CoV-2 E and RdRP genes**

156 The plasmid cDNA encoding for the E and RdRP genes were digested using a pair of
157 restriction sites of the plasmid. Double digestion allowed to isolate the sequence of interest
158 and get linear DNA. The product of digestion was run on a 1% agarose gel with a DNA
159 ladder. The band of interest was excised from the gel and the DNA was purified. To generate
160 positive-sense RNA transcripts, a T7 promoter sequence was added via PCR amplification
161 with the promoter sequence on the forward primer (Supplementary Fig. 6a). The PCR

162 products were verified on an agarose gel (Supplementary Fig. 6b). *In vitro* transcription was
163 done with 2.5 hours incubation, with several rounds of DNase I treatment to remove the DNA
164 template, and the RNA was purified. The RNA was tested by PCR using the RPA primers
165 (also suitable for PCR) to check for traces of DNA impurities (Supplementary Fig. 6c). The
166 concentration of the RNA transcripts was measured using the Qubit, then the RNA was
167 diluted in DEPC-treated water and stored at -80°C with RNase inhibitor. A dilution series
168 was used to measure the analytical sensitivity of the molecular test.

169

170 **Multiplex RT-RPA with complementary detection platforms**

171 *Real-time fluorescence readout:* Amplification and detection of both genes was done using
172 the multi-channel portable reader (Axxin) using the FAM and HEX channels.

173 The 50 µL reactions contained TwistAmp® exo RPA pellets resuspended in 29.5 µL
174 Rehydration Buffer (TwistDX), 2.1 µL of both forward primers (at concentration 10 µM), 2.1
175 µL of both reverse primers (at concentration 10 µM), 0.6 µL of both probes, 1 µL of each
176 corresponding RNA samples (E and RdRP genes), 2.5 µL of reverse transcriptase (at 200
177 U/µL) and 3.9 µL of nuclease-free water. Finally, 2.5 µL of magnesium acetate (at
178 concentration 280 mM) was added to start the reaction. The reactions were incubated at 39°C,
179 with magnetic shaking and the fluorescence was measured in real-time directly from the
180 tubes.

181 *Dipstick readout:* RPA primers for E and RdRP genes were modified (Supplementary Table
182 1) for duplex detection on the dipsticks, which incorporate carbon nanoparticles conjugated
183 to neutravidin. The E gene primers were modified with biotin and digoxigenin for detection
184 on test line (1), whereas the RdRP gene primers were modified with FAM and biotin for
185 detection on test line (2). To eliminate non-specific binding due to dimers forming, the assay
186 was tested without any template (negative controls) with modified primers at concentration

187 10 μ M, 2 μ M, 1 μ M and 0.5 μ M (Supplementary Fig. 4). Eventually, the 50 μ L reactions
188 contained TwistAmp® basic RPA pellets resuspended in 29.5 μ L Rehydration Buffer
189 (TwistDX), 2.1 μ L of both forward primer (at concentration 1 μ M), 2.1 μ L of both reverse
190 primer (at concentration 1 μ M), 1 μ L of each corresponding RNA sample (E and RdRP
191 genes), 2.5 μ L of reverse transcriptase (at 200 U/ μ L) and 5.1 μ L of nuclease-free water.
192 Finally, 2.5 μ L of magnesium acetate (at concentration 280 mM) was added to start the
193 reaction. The reactions were incubated at 37°C in an incubator for 20 minutes, with shaking
194 at 250 rpm. Then, 10 μ L of reaction was mixed in the well of a microplate with 140 μ L of
195 running buffer, and the dipstick was dipped in the well. The test result was read after 10
196 minutes. A photograph of the strips was taken at this time and image analysis was done on
197 Matlab (R2020b).

198 Detection of RNA transcripts spiked in human saliva was done following the protocol for
199 RT-RPA with dipstick readout, yet the 5.1 μ L of nuclease free-water were replaced by human
200 saliva.

201

202 **Calculation of fluorescence thresholds and analytical sensitivity**

203 Two thresholds were calculated for the RT-RPA protocol using the FAM and HEX dyes. The
204 thresholds were computed from eight NTC reactions. The maximum fluorescence values
205 were taken after background subtraction. The average of these values and the standard
206 deviation were calculated. Finally, the thresholds were calculated as followed:

$$207 \textit{Threshold} = \textit{average}(NTC) + 4.785 \times \textit{standard deviation}(NTC)$$

208 The multiplication factor 4.785 corresponds to the 99.9% confidence interval of the t-
209 distribution with seven degrees of freedom, as per the equation 1 for determining limit of
210 blank²⁶. This high confidence interval was chosen to strengthen the specificity of the assay.

211 The analytical sensitivity of the RT-RPA with real-time fluorescence readout was done using
212 these thresholds. Repeats were run five times for a range of RNA inputs: 1, 2.5, 5, 7.5, 10,
213 10^2 and 10^3 (only for the RdRP gene). The fraction of positive reactions (reactions which
214 reached the threshold in less than 20 minutes) was calculated separately for both genes and
215 probit analysis was done on Matlab (R2020b).

216 The EC_{95} was calculated from the probit analysis with its 95% CI. The EC_{95} was defined as
217 the analytical sensitivity of the test.

218

219 **Testing of coronavirus specificity panel**

220 RT-RPA protocol (for both real-time and dipstick readout) was followed to test cross-
221 reactivity of the assay with other coronaviruses, namely SARS-CoV, MERS-CoV, HCoV-
222 NL63, HCoV-OC43 and HCoV-229E. As the RNA concentrations of the stock RNA
223 received from supplier was unknown, the RT-RPA assays were run with 5 μ L of RNA
224 directly from stock. The SARS-CoV-2 RNA sample supplied with the specificity panel was
225 also run with 5 μ L from stock for comparison with the other coronaviruses.

226

227 **Smartphone application**

228 The “CovidApp” smartphone application was developed in Android Studio using java
229 libraries. Screenshots were taken from the emulator using a Galaxy Nexus API 28. The
230 complete code for the Android application is available on request.

231 The application opens onto a homepage where the users can choose between three activities
232 “Test”, “Alerts” or “Map outbreak”. The “Test” activity includes first recording of patient
233 information (patient ID, date of birth, GPS and symptoms). The GPS coordinates are
234 captured in real-time by clicking the button and time and date are also automatically
235 captured. Then, the user can click on the “Take Test Picture” button to get access to the

236 smartphone camera and take a photograph of the lateral flow test. Manual cropping is
237 required to crop onto the result area of the test (where the lines are). Another activity enables
238 image analysis of the cropped image for enhanced visualisation of the test lines and plotting
239 of the test line intensity. Finally, the user can select between the options “three lines”, “two
240 lines”, “one line”, “no line” which records the test result as “positive”, “presumptive
241 positive”, “negative” or “invalid”. If the user test result is “positive” or “presumptive
242 positive”, the user is taken to the “Contacts” page when clicking on the “Next step” button.
243 This will enable to record the contacts of the positive case so then can be later reached by the
244 local contact tracing system. Finally, the activity “Map outbreak” opens to visualise the
245 location of the tested case on the map. In the “Alerts” activity, the information of the patient,
246 with the test result, can be seen.

247

248 **Comparison of four isothermal amplification methods**

249 RT-RPA for visual detection on dipstick was commonly done using a laboratory incubator
250 (New Brunswick™ Innova® 42) at 37°C with shaking at 250 rpm. Other incubation methods
251 were tried and compared including incubation in a water bath at 37°C, incubation on a hand
252 warmer bag (HotHands® air activated) releasing heat at ~36°C after activation and by holding
253 the tube in the hand. The positive and negative reactions using these different incubation
254 methods were all done in parallel with an incubation time of 20 minutes. Then, the reactions
255 were analysed on dipsticks following the dipstick readout protocol.

256

257 **Results**

258 **Gene screening for detection of SARS-CoV-2 by real-time fluorescence**

259 A pair of RPA primers and a fluorescent “exo” probe (Supplementary Table 1) were designed
260 to target four conserved regions of the SARS-CoV-2 genome in the nucleocapsid (N) gene,

261 the envelope (E) gene, the RNA-dependent RNA polymerase (RdRP) gene and the open-
262 reading frame 1a/b (Orf1ab). The RPA assay design was optimised for amplicon size of ~200
263 bp and long primers of ~30 bp. Preliminary BLAST analysis indicated that these pairs and
264 probes specifically detect SARS-CoV-2.

265 A preliminary gene screening aimed to identify the best two primers/probe sets among these
266 four targets, able to achieve rapid and sensitive detection of SARS-CoV-2 in a real-time RPA
267 assay. The gene screening was conducted with cDNA controls rapidly made available by
268 suppliers (Supplementary Fig. 2a). A single fluorescence threshold was used to compare the
269 four targets. All reactions using template, except one (50 copies for the N gene), showed
270 successful amplification of 50, 500 and 5,000 copies with fluorescent signals reaching the
271 threshold in less than 30 minutes (Supplementary Fig. 2b). Then, the average time to
272 threshold was determined for each gene and it was used to compare them (Supplementary
273 Fig. 2c). The two genes with the shortest average time to threshold with 50 copies of cDNA
274 were the E gene, in 14 minutes, and the RdRP gene, in 19 minutes. The Orf1ab gene was
275 slightly slower than RdRP gene, while the N gene showed particularly low sensitivity in the
276 RPA protocol and did not reach the threshold with 50 copies. Eventually, the E and RdRP
277 genes were selected and multiplexed to make an in-house duplex RT-RPA protocol to detect
278 SARS-CoV-2 virus. An analysis of genome variations (determined from 5139 sequenced
279 genomes deposited on <https://www.gisaid.org/>) for the selected primers and probes confirmed
280 that they target conserved regions of the SARS-CoV-2 genome with low variability
281 comprised between 0.1-0.5% (Supplementary Fig. 3).

282

283 **Development of the duplex RT-RPA platforms**

284 The RT-RPA assay was developed with two complementary detection systems (Fig. 1a).

285 First, an optical fluorescent readout similar to qRT-PCR that uses fluorescent probes to

286 monitor real-time amplification of the target was made by multiplexing fluorophores to
287 simultaneously detect the amplicons of the E gene with FAM and RdRP gene with HEX (Fig
288 1.c). The fluorescent probe was designed as a short ~45-50 oligonucleotide sequence,
289 complementary to the target sequence. The fluorescent probe included a fluorophore and a
290 proximal quencher, separated by a tetrahydrofuran (THF) residue. When the fluorescent
291 probe recognised the target sequence, it annealed and was cleaved at the THF site by the
292 exonuclease contained in the exo RPA reaction. As the fluorophore was released from its
293 quencher, a fluorescent signal was produced and recorded on a multi-channel and portable
294 fluorescence reader.

295 A second detection method was developed using dipsticks to detect the amplicons on a
296 nitrocellulose strip using nanoparticle labels. The dipstick-based platform was developed to
297 be as low-cost and minimalist as possible. The primer sequences used were the same as for
298 the real-time fluorescence readout above, but these primers were modified with small
299 molecules to mediate capture of the amplicons on the test lines of the dipstick
300 (Supplementary Table 1). Optimisation of the primer concentration was needed to eliminate
301 non-specific binding on the test lines (Supplementary Fig. 4) attributed to binding of
302 dimerised primers when used in excess ($> 1 \mu\text{M}$). After the amplification was performed,
303 detection of the two amplicons was possible on two distinct test lines: (1) for the E gene, (2)
304 for the RdRP gene and a control line (C) provided confirmation that the test had worked
305 properly (Fig. 1d).

306

307 **Evaluation of the RT-RPA assay with real-time fluorescence detection**

308 The analytical sensitivity was measured for the real-time RT-RPA assay and defined as the
309 concentration of analyte, here synthetic SARS-CoV-2 viral RNA copies per reaction, that can
310 be detected $\geq 95\%$ of the time ($< 5\%$ false negative rate).

311 To determine the analytical sensitivity of the RT-RPA fluorescence readout two thresholds
312 were calculated, to account for the different background fluorescence of the FAM and HEX
313 fluorophores (see Material and methods section). The resulting thresholds were 112 for the E
314 gene (FAM) and 13 for the RdRP gene (HEX). RT-RPA reactions were run for different
315 RNA inputs ranging from 1 copy to 10^5 copies and real-time fluorescence was recorded. The
316 time to threshold was determined for reactions reaching threshold in 20 minutes of
317 amplification (Fig. 2a). The amplification time was fixed at 20 minutes, as the assay was able
318 to detect as little as 1 RNA. To measure the analytical sensitivity of both genes, we calculated
319 the fraction positive to find and plot the EC_{95} (see “Methods” section). The analytical
320 sensitivity was 9.5 RNA copies per reaction (95% CI: 7.0-18) for the E gene and 17 RNA
321 copies per reaction (95% CI: 11-93) for the RdRP gene (Fig. 2b).
322 The specificity of the RT-RPA assay was tested with model samples against common
323 seasonal coronaviruses, namely HCoV-NL63, HCoV-OC63 and HCoV-229E, as their
324 symptoms could be easily confused with COVID-19, and we also tested cross-reactivity with
325 SARS-CoV and MERS-CoV, as they are closely related viruses.
326 No cross-reactivity was observed with the primers/probe set targeting the E gene and the
327 RdRP gene when tested with SARS-CoV- and MERS-CoV and the common colds (Fig. 2c
328 and Fig. 2d). A slight increase in background signal could be observed, although remaining
329 comparable to the non-template control (NTC) reaction and the signal remained below the
330 thresholds.

331

332 **Fig. 2 Evaluation of the sensitivity and specificity of the RT-RPA assay with real-time**
333 **fluorescence detection.**

334

335 **Evaluation of the RT-RPA assay with visual dipstick detection**

336 The analytical sensitivity of the dipstick detection method was approximated by running a
337 range of RNA inputs, from 1 to 10⁵ copies. Six replicates were performed (Supplementary
338 Fig. 5) of which one representative dipstick per RNA concentration is shown in Fig. 3a. The
339 test line intensity analysis was used to quantify test line intensity. Single-copy detection was
340 possible for 2/6 repeats (33%), giving in a positive result, defined as both test lines visible by
341 eye or with image analysis. The probit analysis was performed to determine the analytical
342 sensitivity of the assay using the fraction positive (Fig. 3b). The analytical sensitivity of the
343 dipstick method was 130 (95% CI: 82-500) RNA copies per reaction.
344 The specificity of the dipstick detection method was assessed against the common seasonal
345 coronaviruses, SARS-CoV and MERS-CoV (Fig. 3c). The dipstick showed high specificity
346 for only SARS-CoV-2 viral RNA and no cross-reactivity was seen with the other
347 coronaviruses.

348

349 **Fig. 3 Evaluation of the sensitivity and specificity of the RT-RPA assay with dipstick**
350 **detection.**

351

352 **Exploration of point-of-care testing with the RT-RPA dipstick method**

353 We investigated the potential of the RT-RPA assay for detection of SARS-CoV-2 at the
354 point-of-care with the dipstick readout, a format that could dramatically widen access to
355 testing in decentralised settings.

356 The tests could be read visually by eye. In addition, we developed a smartphone application
357 as a prototype towards a connected-diagnostic dipstick test. The architecture and screenshots
358 of the prototype application are presented in Fig. 4a. The smartphone application proposed
359 allowed input and storage of patient's information, symptoms, capture of geo-location, test
360 lines intensity analysis of the dipstick and a record of the test results. If the test is "Positive"

361 or “Presumptive Positive” the user could insert the names of close contacts for contact tracing
362 purposes. The application also included geographic visualisation of the tested patients to map
363 ‘hotspots’.

364

365 The major advantage of RPA, compared to other approaches such as PCR and LAMP, is its
366 isothermal amplification at $\sim 37^{\circ}\text{C}$. We investigated the potential of different incubation
367 methods which could be more suitable for point-of-care settings. RT-RPA was performed to
368 detect 100 copies of RNA using four incubation approaches: an incubator, a water bath, a
369 disposable hand warmer bag and simply holding the tube in our hands. Incubators and water
370 baths are often found in well-equipped laboratories, but we also tried using a low-cost hand
371 warmer bag (based on an exothermic reaction shown to deliver a constant temperature of
372 $\sim 36\text{-}37^{\circ}\text{C}$ for several hours²⁷) and holding the tube in one hand (using body temperature
373 $\sim 37^{\circ}\text{C}$) to show inexpensive and equipment-free alternatives. The results are shown in Fig.
374 4b. While amplification in the incubator seemed to show the best results with two test lines
375 visible on the dipstick, two test lines were also visible for the reaction incubated in the water
376 bath, although slightly fainter. The reactions incubated on a hand warmer bag and handheld
377 appeared less sensitive, showing only a signal on test line (1). However, we proved that very
378 simple methods could be successfully used to amplify SARS-CoV-2 RNA via RT-RPA and
379 visual dipstick detection.

380

381 Finally, preliminary analysis was performed to assess the potential of the dipstick test to be
382 compatible with mock clinical samples, using human saliva with spiked RNA transcripts to
383 mimic mouth swabs (Fig. 4c). Saliva is an easy specimen for self-collection that has been
384 FDA-approved for molecular testing of COVID-19²⁸. The E gene was clearly detectable on
385 test line (1) with ≥ 1 RNA copy per reaction, and a faint signal was seen on test line (2) for

386 the RdRP gene with 1 and 100 RNA copies per reaction. Two strong test lines were visible
387 for 10⁵ copies per reaction. Therefore, the findings of this small study suggest that conducting
388 the assay in saliva compared to buffer did not have a substantial impact on the assay
389 sensitivity.

390

391 **Fig. 4 Exploration of point-of-care testing with the RT-RPA dipstick method.**

392 **Discussion**

393 Herein we report the development and evaluation of a rapid (25-35 minutes), multiplexed
394 molecular diagnostic for SARS-CoV-2 by RT-RPA. The test was presented with two
395 complementary detection methods: real-time fluorescence using a portable reader and visual
396 dipstick readout on a low-cost nitrocellulose strip. The test showed high sensitivity and high
397 specificity for both readouts. The detection method by dipstick was further investigated for
398 point-of-care and decentralised testing using different incubation methods and a smartphone
399 application to capture, analyse and connect test results.

400

401 The development of the multiplex isothermal RT-RPA assay started by selecting two optimal
402 targets, in the E gene and RdRP gene, for rapid and ultrasensitive detection of SARS-CoV-2.
403 Detecting several targets in a multiplex test was done to increase the robustness of the assay,
404 as a common strategy also seen with PCR protocols (Supplementary Fig. 1).

405

406 Isothermal fluorescence readers are usually available in centralised laboratories; however,
407 they are not necessarily found in decentralised laboratories and low-resource settings. For this
408 reason, we developed a second readout format, using a dipstick. Dipsticks are portable, cost-
409 effective and user-friendly tools that can detect RPA amplicons with minimal equipment and
410 the test result can be seen with the naked eye. Here only a tube or microplate to mix the RT-

411 RPA reaction with the buffer and a pipette to apply the mix on the dipstick was needed.

412 Giving the option of two alternative readouts with their own advantages aimed to make

413 molecular testing more widely accessible and suitable for decentralised testing.

414

415 The amplification time for the RT-RPA assay was set to run for 20 minutes as it was enough

416 to achieve single-copy detection of the E gene with real-time fluorescence and visual dipstick

417 readouts, showing the ultrasensitive potential of the test. The analytical sensitivities for the E

418 and RdRP genes comparable with those reported by Charité for its qRT-PCR assay which

419 were 5.2 copy per reaction (95% CI: 3.7-9.6) for the E gene and 3.8 copy per reaction (95%

420 CI: 2.7-7.6) for the RdRP gene, in comparison to 9.5 RNA copies per reaction (95% CI: 7.0-

421 18) for the E gene and 17 RNA copies per reaction (95% CI: 11-93) for the RdRP gene

422 reported herein. It was necessary to achieve high sensitivity for the molecular test to detect

423 viral loads that are clinically relevant for the disease, here COVID-19. The RT-RPA assay

424 was shown to be highly specific to SARS-CoV-2, with no observed cross-reactivity with the

425 closely related coronaviruses tested, such as SARS-CoV, MERS-CoV, HCoV-NL63, HCoV-

426 OC43 and HCoV-229E. This high specificity was demonstrated for both detection methods

427 and reduces the risk of false positives with closely related viruses.

428

429 The prototype smartphone application was proposed as a powerful tool for data capture,

430 analysis and visualisation when testing in decentralised settings. Smartphones are widely

431 accessible, easy-to-use and can act as a substitute to sophisticated laboratory equipment as

432 they integrate a high-resolution camera, large data storage space, real-time location and

433 connectivity. Altogether, these features make health-related smartphone applications

434 attractive accessories to elevate point-of-care diagnostic tests. Moreover, the use of

435 inexpensive methods for incubation at 37°C of the RT-RPA reaction for detection on

436 dipsticks, especially with a hand warmer bag (~ \$0.5) and using body temperature,
437 emphasised the simplicity of the assay.

438

439 **Conclusion**

440 To close, we have developed an ultrasensitive and specific diagnostic for SARS-CoV-2 viral
441 RNA using isothermal RPA technology, and proposed two different detection methods, both
442 showing high accuracy. While real-time fluorescence detection developed here offers more
443 sensitivity and faster results (10 minutes faster than dipstick method), the proposed detection
444 on dipsticks appeared as the best method for decentralised testing. Having an alternative to
445 qRT-PCR that is comparable in performance, but with a shorter time-to-result, using different
446 supply chains, requiring less equipment and non-extensive laboratory experience could help
447 to alleviate the pressure on healthcare systems and curb the COVID-19 pandemic worldwide.

448

449 Further test development will include clinical validation of the RT-RPA assay with clinical
450 samples with cross-validation of the developed assay with qRT-PCR results to determine the
451 clinical sensitivity and specificity of the test. Preliminary analysis of the dipstick readout with
452 human saliva showed no cross-reaction between mock clinical samples and the RT-RPA
453 assay, hence saliva appeared as a good specimen candidate for a non-invasive test using
454 mouth swab. In future, the adaption of multiplexed gene analysis to detect the S gene could
455 help to track the proportion of new variants of concern and the impact of COVID-19
456 vaccinations.

457

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466 **References**

- 467 1. Wu, F. *et al.* A new coronavirus associated with human respiratory disease in China.
468 *Nature* (2020). doi:10.1038/s41586-020-2008-3
- 469 2. Gorbalenya, A. E. *et al.* The species Severe acute respiratory syndrome-related
470 coronavirus: classifying 2019-nCoV and naming it SARS-CoV-2. *Nat. Microbiol.*
471 (2020). doi:10.1038/s41564-020-0695-z
- 472 3. World Health Organization. WHO Director-General's remarks at the media briefing on
473 2019-nCoV on 11 February 2020. Available at:
474 [https://www.who.int/dg/speeches/detail/who-director-general-s-remarks-at-the-media-](https://www.who.int/dg/speeches/detail/who-director-general-s-remarks-at-the-media-briefing-on-2019-ncov-on-11-february-2020)
475 [briefing-on-2019-ncov-on-11-february-2020](https://www.who.int/dg/speeches/detail/who-director-general-s-remarks-at-the-media-briefing-on-2019-ncov-on-11-february-2020). (Accessed: 25th August 2020)
- 476 4. World Health Organization. WHO Coronavirus Disease (COVID-19) Dashboard.
477 Available at:
478 https://covid19.who.int/?gclid=Cj0KCQjw7ZL6BRCmARIsAH6XFDI6_x2M3aCcbu
479 [kdjbY-VVqN2TONEsTcjuRMEu7A6nqss4ydtGlcYFsaAj8lEALw_wcB](https://covid19.who.int/?gclid=Cj0KCQjw7ZL6BRCmARIsAH6XFDI6_x2M3aCcbu). (Accessed:
480 8th January 2021)
- 481 5. World Health Organization. WHO Director-General's opening remarks at the media
482 briefing on COVID-19 - 13 April 2020. (2020). Available at:
483 [https://www.who.int/dg/speeches/detail/who-director-general-s-opening-remarks-at-](https://www.who.int/dg/speeches/detail/who-director-general-s-opening-remarks-at-the-media-briefing-on-covid-19--13-april-2020)
484 [the-media-briefing-on-covid-19--13-april-2020](https://www.who.int/dg/speeches/detail/who-director-general-s-opening-remarks-at-the-media-briefing-on-covid-19--13-april-2020). (Accessed: 29th June 2020)
- 485 6. Hu, Z. *et al.* Clinical characteristics of 24 asymptomatic infections with COVID-19

- 486 screened among close contacts in Nanjing. *China. Sci China Life Sci* (2020).
487 doi:10.1007/s11427-020-1661-4
- 488 7. Nishiura, H., Linton, N. M. & Akhmetzhanov, A. R. Serial interval of novel
489 coronavirus (COVID-19) infections. *Int. J. Infect. Dis.* (2020).
490 doi:10.1016/j.ijid.2020.02.060
- 491 8. World Health Organization. Laboratory testing of 2019 novel coronavirus (2019-
492 nCoV) in suspected human cases: interim guidance, 17 January 2020. Available at:
493 [https://www.who.int/publications/i/item/laboratory-testing-of-2019-novel-coronavirus-](https://www.who.int/publications/i/item/laboratory-testing-of-2019-novel-coronavirus-(-2019-ncov)-in-suspected-human-cases-interim-guidance-17-january-2020)
494 [\(-2019-ncov\)-in-suspected-human-cases-interim-guidance-17-january-2020](https://www.who.int/publications/i/item/laboratory-testing-of-2019-novel-coronavirus-(-2019-ncov)-in-suspected-human-cases-interim-guidance-17-january-2020).
495 (Accessed: 25th August 2020)
- 496 9. National Institute for Viral Disease Control and Prevention. 新型冠状病毒核酸检测
497 引物和探针序列 (Specific primers and probes for detection 2019 novel coronavirus).
498 Available at: http://ivdc.chinacdc.cn/kyjz/202001/t20200121_211337.html. (Accessed:
499 25th August 2020)
- 500 10. Corman, V. M. *et al.* Detection of 2019 novel coronavirus (2019-nCoV) by real-time
501 RT-PCR. *Eurosurveillance* (2020). doi:10.2807/1560-7917.ES.2020.25.3.2000045
- 502 11. Centers for Disease Control and Prevention. *2019-Novel Coronavirus (2019-nCoV)*
503 *Real-time rRT-PCR Panel Primers and Probes.* (2020).
- 504 12. FIND. Testing is our first line of defence. (2020). Available at:
505 <https://www.finddx.org/testing-matters/>. (Accessed: 29th June 2020)
- 506 13. Vandenberg, O., Martiny, D., Rochas, O., van Belkum, A. & Kozlakidis, Z.
507 Considerations for diagnostic COVID-19 tests. *Nat. Rev. Microbiol.* (2020).
508 doi:10.1038/s41579-020-00461-z
- 509 14. GOV.UK. UK reaches 200,000 coronavirus testing capacity target a day early -

- 510 GOV.UK. Available at: <https://www.gov.uk/government/news/uk-reaches-200000->
511 [coronavirus-testing-capacity-target-a-day-early](https://www.gov.uk/government/news/uk-reaches-200000-coronavirus-testing-capacity-target-a-day-early). (Accessed: 10th September 2020)
- 512 15. Sheridan, C. COVID-19 spurs wave of innovative diagnostics. *Nat. Biotechnol.*
513 (2020). doi:10.1038/s41587-020-0597-x
- 514 16. TwistDx. Recombinase Polymerase Amplification (RPA). Available at:
515 <https://www.twistdx.co.uk/en/rpa>. (Accessed: 29th June 2020)
- 516 17. Daher, R. K., Stewart, G., Boissinot, M. & Bergeron, M. G. Recombinase Polymerase
517 Amplification for Diagnostic Applications. *Clin. Chem.* (2016).
518 doi:10.1373/clinchem.2015.245829
- 519 18. Mori, Y. & Notomi, T. Loop-mediated isothermal amplification (LAMP): a rapid,
520 accurate, and cost-effective diagnostic method for infectious diseases. *J. Infect.*
521 *Chemother.* (2009). doi:<https://doi.org/10.1007/s10156-009-0669-9>
- 522 19. U.S. Food and Drug Administration (FDA). Coronavirus (COVID-19) Update: FDA
523 Authorizes First COVID-19 Test for Self-Testing at Home. (2020). Available at:
524 [https://www.fda.gov/news-events/press-announcements/coronavirus-covid-19-update-](https://www.fda.gov/news-events/press-announcements/coronavirus-covid-19-update-fda-authorizes-first-covid-19-test-self-testing-home)
525 [fda-authorizes-first-covid-19-test-self-testing-home](https://www.fda.gov/news-events/press-announcements/coronavirus-covid-19-update-fda-authorizes-first-covid-19-test-self-testing-home). (Accessed: 18th November 2020)
- 526 20. Xia, S. & Chen, X. Single-copy sensitive, field-deployable, and simultaneous dual-
527 gene detection of SARS-CoV-2 RNA via modified RT-RPA. *Cell Discov.* (2020).
528 doi:10.1038/s41421-020-0175-x
- 529 21. Behrmann, O. *et al.* Rapid detection of SARS-CoV-2 by low volume real-time single
530 tube reverse transcription recombinase polymerase amplification using an exo probe
531 with an internally linked quencher (exo-IQ). (2020).
532 doi:10.1093/clinchem/hvaa116/5834714
- 533 22. Miller, B. S. *et al.* Spin-enhanced nanodiamond biosensing for ultrasensitive
534 diagnostics. *Nature* (2020). doi:10.1038/s41586-020-2917-1

- 535 23. Wood, C. S. *et al.* Taking Mobile Health Connected Infectious Disease Diagnostics to
536 the Field. *Nature* (2019). doi:10.1038/s41586-019-0956-2
- 537 24. Brangel, P. *et al.* A Serological Point-of-Care Test for the Detection of IgG Antibodies
538 against Ebola Virus in Human Survivors. *ACS Nano* (2018).
539 doi:10.1021/acsnano.7b07021
- 540 25. Budd, J. *et al.* Digital technologies in the public-health response to COVID-19. *Nat.*
541 *Med.* (2020). doi:10.1038/s41591-020-1011-4
- 542 26. Holstein, C. A., Griffin, M., Hong, J. & Sampson, P. D. Statistical Method for
543 Determining and Comparing Limits of Detection of Bioassays. *Anal. Chem* (2015).
544 doi:10.1021/acs.analchem.5b02082
- 545 27. Wang, L. What's inside disposable hand warmers? *Chemical and Engineering News*
546 (2010). Available at: <https://cen.acs.org/articles/88/i4/Hand-Warmers.html>.
- 547 28. U.S. Food and Drug Administration (FDA). Coronavirus (COVID-19) Update: FDA
548 Authorizes First Diagnostic Test Using At-Home Collection of Saliva Specimens.
549 Available at: [https://www.fda.gov/news-events/press-announcements/coronavirus-](https://www.fda.gov/news-events/press-announcements/coronavirus-covid-19-update-fda-authorizes-first-diagnostic-test-using-home-collection-saliva)
550 [covid-19-update-fda-authorizes-first-diagnostic-test-using-home-collection-saliva.](https://www.fda.gov/news-events/press-announcements/coronavirus-covid-19-update-fda-authorizes-first-diagnostic-test-using-home-collection-saliva)
551 (Accessed: 15th October 2020)

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558

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571

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575 Da Huang: Conceptualisation, methodology, formal analysis, visualisation, project
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578 Rachel A. McKendry: Conceptualisation, funding acquisition, supervision, writing – review
579 & editing

580

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582

583 **Supplementary information**

584 File: Supplementary Table and Figures.docx

1 **Figure captions**

2

3 **Harnessing recombinase polymerase amplification for** 4 **rapid detection of SARS-CoV-2 in resource-limited** 5 **settings**

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17 **Fig. 1 Schematic representation of the rapid and multiplex RT-RPA assay with real-**
18 **time fluorescence and dipstick detection.**

19 **a** One-pot RT-RPA assay including reverse transcription of the viral RNA and amplification
20 by RPA at constant temperature (37-39°C). **b** Sequences of the primers/probe sets used for
21 SARS-CoV-2 E gene and RdRP gene in the multiplex RT-RPA assay with real-time
22 detection (blue) and sequences of the modified primers used for the multiplex dipstick
23 detection (orange). **c** Real-time fluorescence detection by exonuclease cleavage of the probes
24 for E gene and RdRP gene at their THF residue. **d** Design of the dipstick for multiplexed
25 detection of the E gene and the RdRP gene.

26

27 **Fig. 2 Evaluation of the sensitivity and specificity of the RT-RPA assay with real-time**
28 **fluorescence detection.**

29 **a** Time to threshold for positive RT-RPA reactions with real-time fluorescence detection. The
30 dots represent individual values for the positive reactions reaching threshold for the E gene
31 (in green) and for the RdRP gene (in orange). The bars represent the average time to
32 threshold for the positive reactions and the error bars represent the standard deviation. Each
33 RNA concentration was run in five replicates (N=5), only the positive reactions are
34 represented. **b** Probit analysis for the E gene (left, green) and RdRP gene (right, orange) with
35 their 95% confidence interval (CI). The fraction positive was determined from the RT-RPA
36 reactions in **a** and the probit analysis was done to find the effective concentration at 95%
37 (EC₉₅) for both genes. **c** Validation of the specificity of the E gene primers/probe set against
38 SARS-CoV, MERS-CoV (top) and the seasonal coronaviruses (bottom). **d** Validation of the
39 specificity of the RdRP gene primers/probe set against SARS-CoV, MERS-CoV (top) and the
40 seasonal coronaviruses (bottom). NTC: non-template control.

41

42 **Fig. 3 Evaluation of the sensitivity and specificity of the RT-RPA assay with dipstick**
43 **detection.**

44 **a** Evaluation of the sensitivity of the RT-RPA with multiplex dipstick detection. Captures of
45 the dipsticks run with a range of RNA inputs are shown with the associated test line intensity
46 analysis. Dipsticks were annotated (- -) if no test line was visible (“Negative”), (+ -) or (- +)
47 if only one test line was visible (“Presumptive positive”) and (+ +) if both test lines (1) and
48 (2) were visible (“Positive”). One representative dipstick capture is shown here. **b** Probit
49 analysis and determination of the EC₉₅ for the dipstick detection method (taking both genes
50 into account) with the 95% confidence interval (CI). The fraction positive was determined
51 from six replicates (N=6) RT-RPA reactions. **c** Specificity of the dipstick detection method
52 against SARS-CoV, MERS-CoV (left) and the seasonal coronaviruses (right). Photographs of
53 the dipsticks are shown (top) with the associated test line intensity analysis (bottom). NTC:
54 non-template control.

56 **Fig. 4 Exploration of point-of-care testing with the RT-RPA dipstick method.**

57 **a** Architecture of the prototype smartphone application “CovidApp”. The design of the
58 smartphone application is represented along with screenshots of the different activities of the
59 application. The main activities, including “Homepage”, “Test”, “Contact”, “Alerts” and
60 “Map Outbreak” are described. **b** Comparison of incubation of four methods for RT-RPA at
61 ~37°C with dipstick readout, including incubation using a traditional laboratory incubator
62 with shaking, water bath, hand warmer bag and handheld (using body temperature)
63 (photographs on the top). The lateral flow test captures are shown (middle) with the
64 associated test line intensity analysis (bottom). **c** Detection of RT-RPA reactions with mock
65 clinical samples (saliva spiked with RNA). Photographs of the lateral flow test captures are
66 shown (top) with the associated test line intensity analysis (bottom). **b** and **c** dipsticks were

- 67 annotated (- -) if no test line was visible (“Negative”), (+ -) or (- +) if only one test line was
68 visible (“Presumptive positive”) and (+ +) if both test lines (1) and (2) were visible
69 (“Positive”). NTC: non-template control; PC: positive control (100 copies RNA/reaction).

a RT-RPA (37-39°C)

Viral RNA

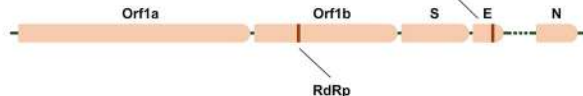


cDNA amplicons

Real-time
Detection

Forward_E_gene: 5'-TTACACTAGCCATCTTACTGCGCTCGAT-3'

Reverse_E_gene: 5'-GCTAAAATTAAGTCCAAACAGAAAAACT-3'

exo_probe_E_gene: 5'-AGACCAGAAGATGACGAACCTAGAAGAAT[**FAM**][**THF**][**BHQ-1**][GATTTTAAACAGAG-3'Multiplex
Dipstick
DetectionForward_E_gene_modified: 5'-[**DIG**][TTACACTAGCCATCTTACTGCGCTCGAT-3'Reverse_E_gene_modified: 5'-[**Biotin**][GCTAAAATTAAGTCCAAACAGAAAAACT-3'Real-time
Detection

Forward_RdRp_gene: 5'-GAGTGTGCTCAAGTATTGAGTGAATGGTC-3'

Reverse_RdRp_gene: 5'-CTGTGTTGTAATTCGGACATCACTATCG-3'

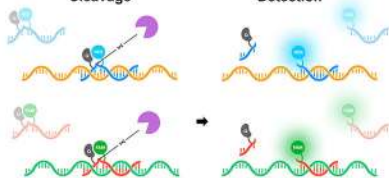
exo_probe_RdRp_gene: 5'-TTACCATCAGTAGATAAAAAGTCATTAACA[**BEX**][**THF**][**BHQ-2**][GCCGTGACAGCTTA-3'Multiplex
Dipstick
DetectionForward_RdRp_gene_modified: 5'-[**Biotin**][GAGTGTGCTCAAGTATTGAGTGAATGGTC-3'Reverse_RdRp_gene_modified: 5'-[**FAM**][CTGTGTTGTAATTCGGACATCACTATCG-3'

5 min

15 min

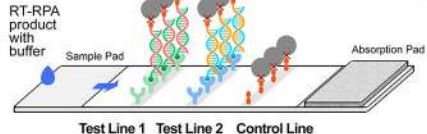
25 min

35 min

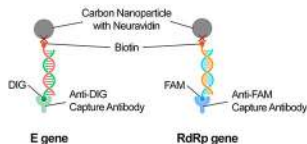
Real-time detection
with portable fluorescence readerDipstick detection
(enhancement with smartphone)**c** Exonuclease
CleavageFluorescence
DetectionE gene probe
with FAMRdRp gene probe
with HEX

Exonuclease

Quencher

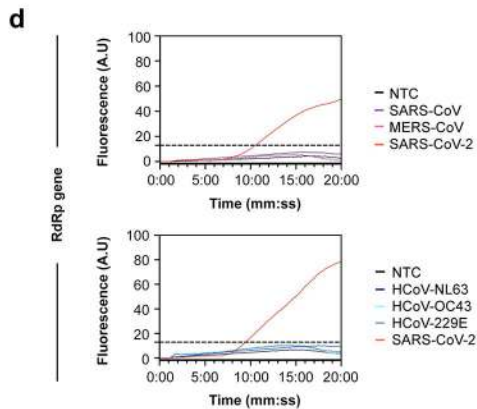
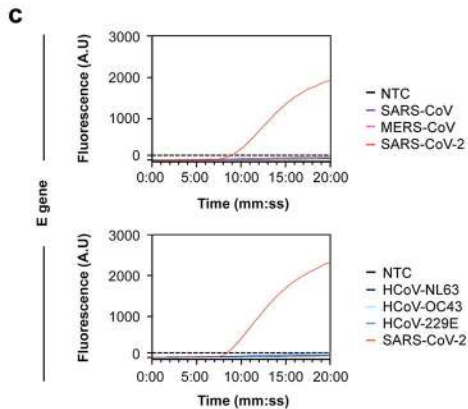
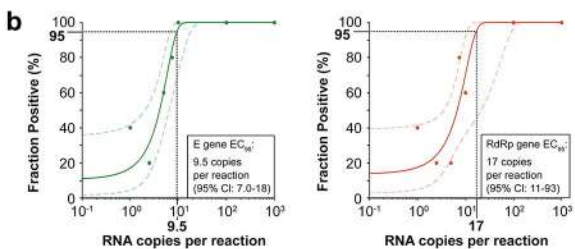
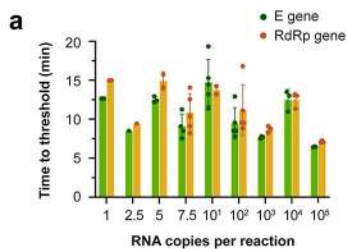
THF
Residue**d**

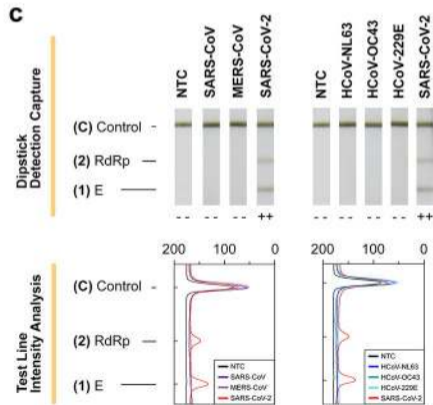
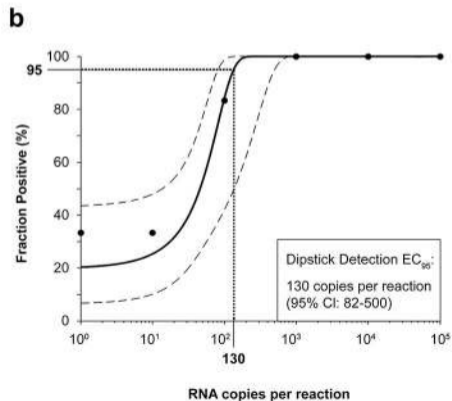
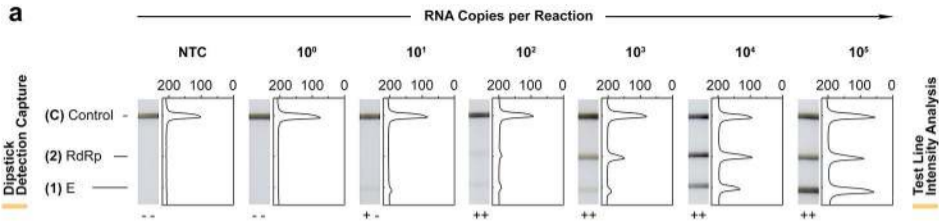
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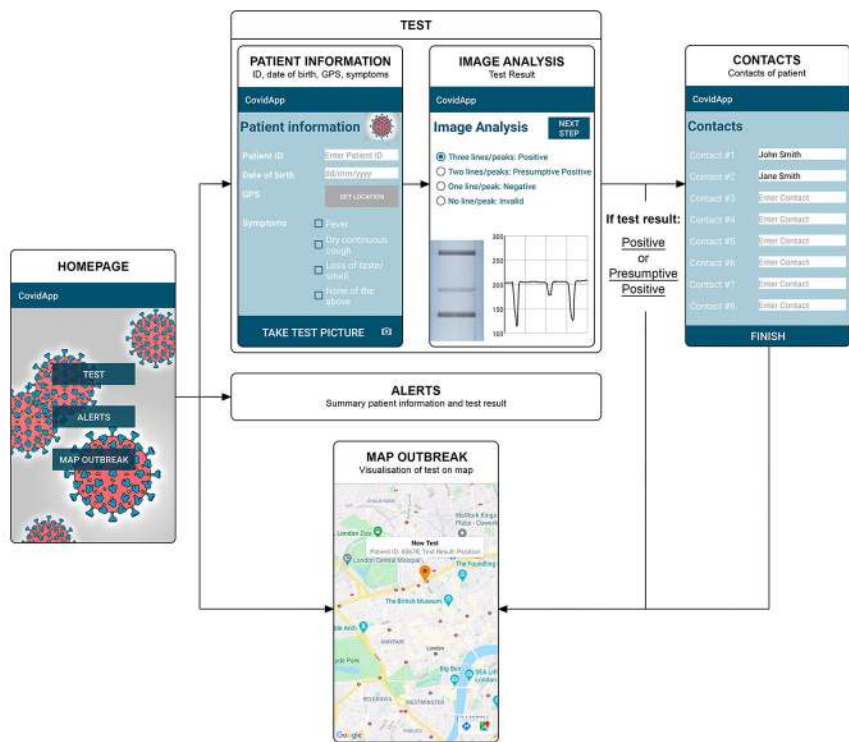
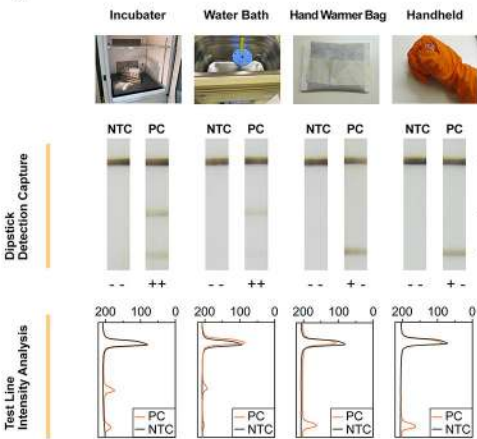


E gene

RdRp gene





a**b****c**