

SUPPLEMENTARY INFORMATION

Massive and rapid COVID-19 testing is feasible by extraction-free SARS-CoV-2 RT-PCR

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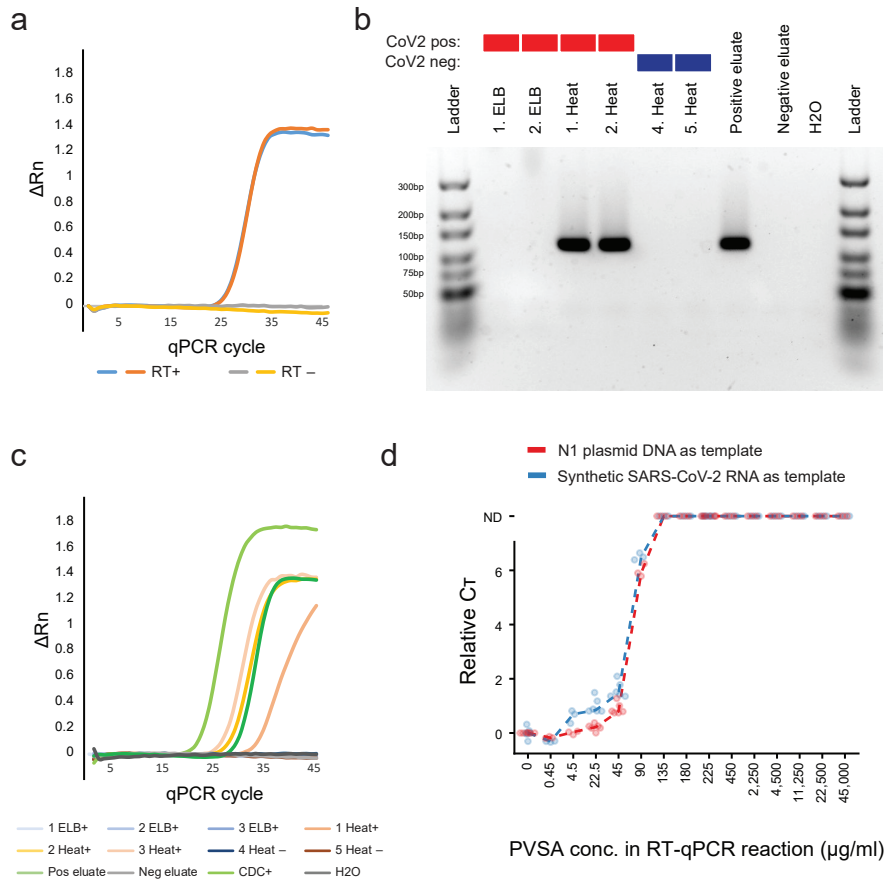
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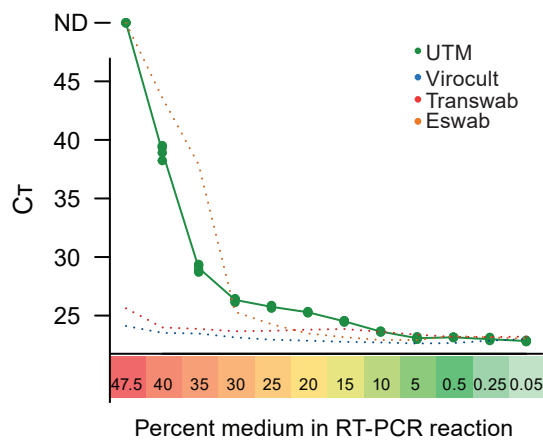
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Supplementary figure 1



Supplementary figure 1. (a) Amplification plots showing normalized reporter value (ΔR_n , linear scale) as a function of qPCR cycle for two-step RT and qPCR performed on synthetic full-genome SARS-CoV-2 RNA (SKU102024-MN908947.3, Twist Biosciences) including (RT+) or excluding (RT-) the reverse transcriptase. (b) Agarose gel electrophoresis of RT-PCR products for reactions described in Fig.2e, but in this case using primers and probes for gene E. (c) Amplification plots showing normalized reporter value (ΔR_n , linear scale) as a function of qPCR cycle for the samples described in Fig. 2e-f, but using two-step RT and qPCR instead of the single-step TaqPath RT-qPCR (Methods). (d) Relative cycle threshold (C_T) values of SARS-CoV-2 RT-PCR using the N1 primer-probe set and a dilution series of the chemical RNase inhibitor polyvinylsulfonic acid (PVSA). To test the inhibition of PVSA in RT and PCR reactions separately, we used either synthetic full-genome SARS-CoV-2 RNA (blue dots and line) or plasmid DNA (CDC positive control) (red dots and line) as template in the RT-PCR reaction. C_T values are shown relative to reactions without PVSA (conc.= 0). The lines indicate the median of triplicate measurements.

Supplementary figure 2



Supplementary figure 2. C_T values from RT-PCR performed on dilution series of Universal Transport Medium (UTM, Copan) using 50,000 spiked copies of synthetic full-genome SARS-CoV-2 RNA and the N1 primer-probe set. Lines represent the mean and the values of individually replicates ($n=4$) are shown as dots. Mean values for Virocult, Transwab and Eswab are shown as dashed lines, included for comparison.

Supplementary table 2

Supplementary table 2. Limit-of-detection data. C_T values for a COVID-19 positive clinical nasopharyngeal swab sample diluted 1:100, 1:1000, 1:10,000 or 1:100,000 in PBS and quantified using cobas 6800 and TaqPath SARS-CoV-2 RT-PCR on eluted RNA.

Sample dilution	Replicate	cobas 6800	RT-PCR (eluate)	cobas 6800	RT-PCR (eluate)
		E	E	ORF1	RdRP
1:100	1	35.16	34.98	35.14	33.18
	2	34.97	ND	34.95	33.89
	3	35.26	32.86	34.60	32.92
1:1000	1	37.35	36.53	36.32	33.90
	2	36.25	35.36	36.06	34.55
	3	36.68	ND	35.88	35.36
1:10,000	1	37.91	35.31	37.05	35.55
	2	38.71	ND	ND	ND
	3	39.09	ND	ND	ND
1:100,000	1	ND	ND	ND	ND
	2	ND	ND	ND	ND
	3	ND	ND	ND	ND

Supplementary table 3

Supplementary table 3. Generic buffers evaluated for SARS-CoV-2 hid-RT-PCR.

Buffer	Composition	Measured pH	Company catalog number (if available)
PVSA in H ₂ O pH6.5	Polyvinylsulfonic acid in H ₂ O (50µg/ml)	6.5	Sigma 278424, Ambion AM9932
TE + 4% sucrose pH7.0	TE (10 mM Tris; 0.1mM EDTA) + 4% sucrose	7.0	Invitrogen AM9858
Tris buffer 10mM pH7.0	10mM Tris buffer	7.0	
PVSA in Tris buffer pH7.0	PVSA in 10mM Tris buffer (50µg/ml PVSA)	7.0	
Tris-HCl 5mM pH7.4	Tris-HCl 5mM	7.4	
TE pH7.0	TE (10 mM Tris + 0.1mM EDTA)	7.0	Invitrogen AM9858
PBS 10mM pH7.5	10mM PBS	7.5	Invitrogen AM9624
Sodium citrate 1mM pH7.6	1mM Sodium citrate	7.6	
H ₂ O	H ₂ O	6.4	Ambion AM9932
PVSA in 1mM Sodium citrate pH7.5	PVSA in 1mM Sodium citrate (50µg/ml PVSA)	7.5	
TE + 4% sucrose pH8.0	TE (10 mM Tris; 0.1mM EDTA) + 4% sucrose)	8.0	Invitrogen AM9858
TE pH8.1	TE (10 mM Tris 0.1mM EDTA)	8.1	Invitrogen AM9858
HBSS pH8.4	HBSS (Hanks' Balanced Salt solution)	8.4	Gibco 14025-092
PBS 100mM pH7.7	100mM PBS	7.7	Invitrogen AM9624
Saline 0.9% pH5.8	0.9% NaCl in 1L water	5.8	
DMEM pH7.9	DMEM (Dulbecco's Modified Eagle Medium)	7.9	Sigma D5796
Citrate buffer pH6.2	Citrate buffer (0.04M Sodium citrate dihydrate; 0.06M Citric acid)	6.2	