

Supplemental Information

Loss of detection of sgN precedes viral abridged replication in COVID19-affected patients – a target for SARS-CoV-2 propagation

Veronica Ferrucci^{1,2,14}, Pasqualino de Antonellis^{1,2,14}, Fabrizio Quarantelli², Fatemeh Asadzadeh^{1,2}, Francesca Bibbò^{1,2}, Roberto Siciliano¹, Carmen Sorice^{1,2}, Ida Pisano¹, Barbara Izzo^{1,2}, Carmela Di Domenico¹, Angelo Boccia¹, Maria Vargas³, Biancamaria Pierri⁴, Maurizio Viscardi⁴, Sergio Brandi⁴, Giovanna Fusco⁴, Pellegrino Cerino⁴, Livia De Pietro⁵, Ciro Furfaro⁶, Leonardo Antonio Napolitano⁶, Giovanni Paolella^{1,2}, Lidia Festa^{7,8}, Stefania Marzinotto⁹, Maria Concetta Conte⁷, Ivan Gentile⁸, Giuseppe Servillo³, Francesco Curcio⁹, Tiziana De Cristofaro^{10,11}, Francesco Broccolo¹², Ettore Capoluongo^{1,2,13} and Massimo Zollo^{1,2,13*}

1 CEINGE Biotecnologie Avanzate, Naples, 80145, Italy.

2 Dipartimento di Medicina Molecolare e Biotecnologie Mediche (DMMBM), 'Federico II' University of Naples, Naples, 80131, Italy.

3 Department of Neurosciences, Reproductive Sciences and Odontostomatology, University of Naples Federico II, Naples, Italy.

4 Istituto Zooprofilattico Sperimentale del Mezzogiorno, Naples, 80055, Italy.

5 U.O.C. Pneumologia, Ospedale Boscotrecase- Asl Napoli3 Sud, 80042 Boscotrecase- Naples, Italy.

6 Dipartimento di Medicina di Laboratorio e Anatomia Patologica, Asl Napoli 3 Sud 80035, Naples, Italy

7 Unità Speciale di Continuità Assistenziale (USCA), Asl Benevento, 82100, Italy.

8 Dipartimento di Medicina Clinica e Chirurgia, AOU Federico II, Naples, Italy

9 Dipartimento di Medicina di Laboratorio, Università degli Studi di Udine, 33100, Udine, Italy

10 IEOS-Institute of Experimental Endocrinology and Oncology G. Salvatore, National Research Council, Naples, 80131, Italy.

11 BioMol Laboratories srl, Naples, 80146, Italy.

12 Laboratory of Molecular Microbiology & Virology, School of Medicine, University of Milano-Bicocca, Monza, 20900, Italy.

13 DAI Medicina di Laboratorio e Trasfusionale, AOU Azienda Ospedaliera 'Federico II', Naples, 80131, Italy.

14 These authors contributed equally.

*Corresponding author. Prof. Massimo Zollo. Email: massimo.zollo@unina.it

***Corresponding author.** Email: massimo.zollo@unina.it

Supplementary Figures

Supplementary Figure S1

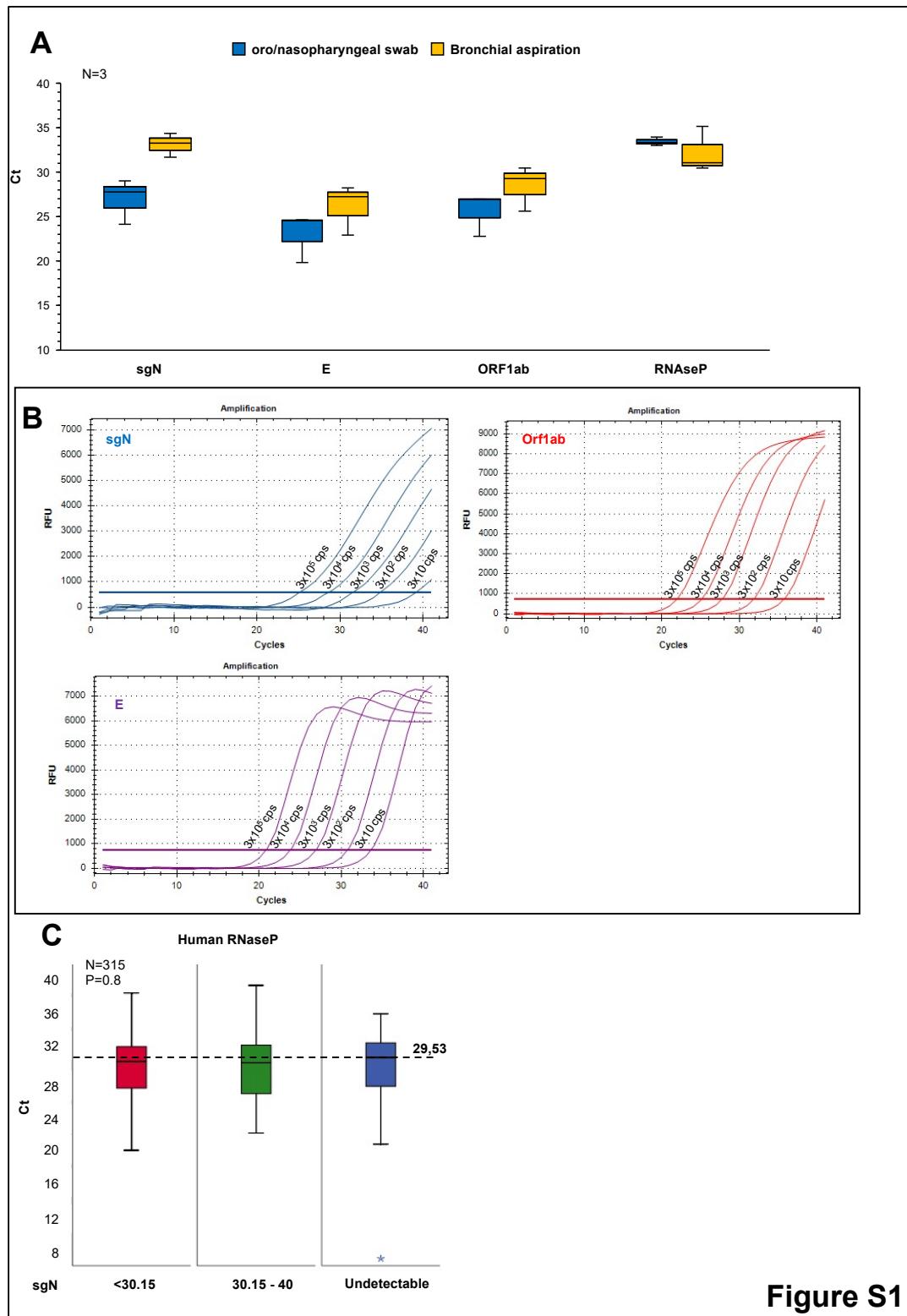


Figure S1

Supplementary Figure S1. SARS-CoV-2 Viral3 kit to detect sgN. Related to Figures 1. (A)

Detection of viral sgN, gene E, gene ORF1ab and human RNase P gene using the SARS-CoV-2 Viral3 kit for oro/nasopharyngeal swabs (orange) and bronchial aspirate samples (orange) in 3 hospitalized COVID19-positive patients. Mean Cq values are shown. **(B)** qPCR assays for the sensitivity of the SARS-CoV-2 Viral3 kit. Human 2019-nCoV strain 2019-nCoV/Italy INMI1 RNA was used as reference to test the sensitivity. Serial dilutions (from 3×10^5 to 3×10^0) of Human 2019-nCoV strain 2019-nCoV/Italy INMI1 RNA were used. cps, standard copy particles. **(C)** Samples obtained from oro/nasopharyngeal swabs from COVID19-positive patients ($N = 315$) were stratified into three groups according to the median Cq values of sgN (sgN Cq median = 33.51). The first group consisted of those samples where Cq values for sgN were below the median value (i.e., Cq < 30.51; 99 samples, in red). The second group of samples were characterized by Cq values of sgN ranged from the Cq median value (30.51) to 40 (96 samples, in green). Third group comprised samples in which sgN was not detectable (i.e., Cq > 40; 120 samples, in blue). One-way analysis of variance (ANOVA) was used through IBM SPSS Statistics to determine the cut-off for sgN detection. SgN was detected in these samples independent of human RNaseP detection ($P < 0.8$).

Supplementary Figure S2.

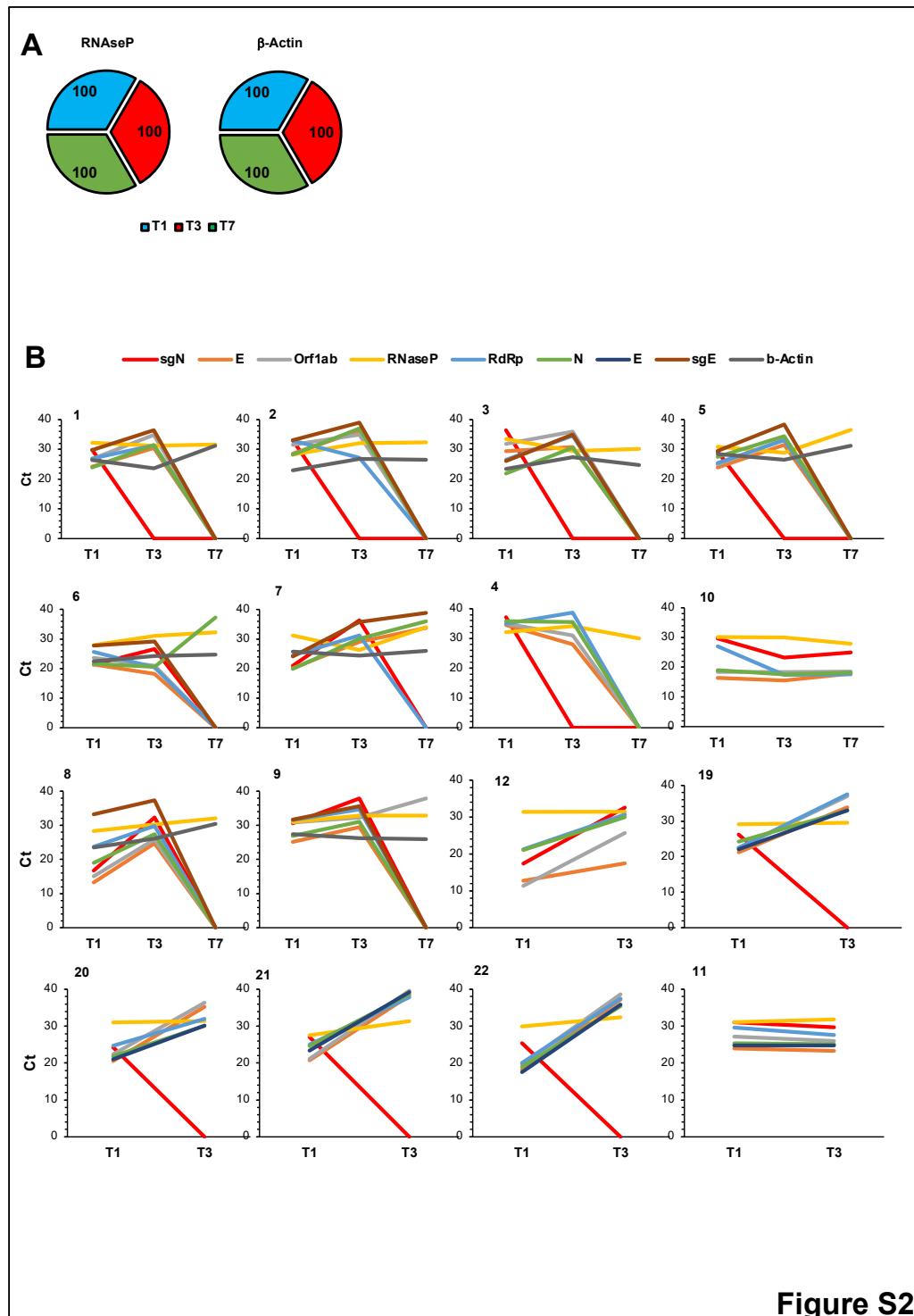
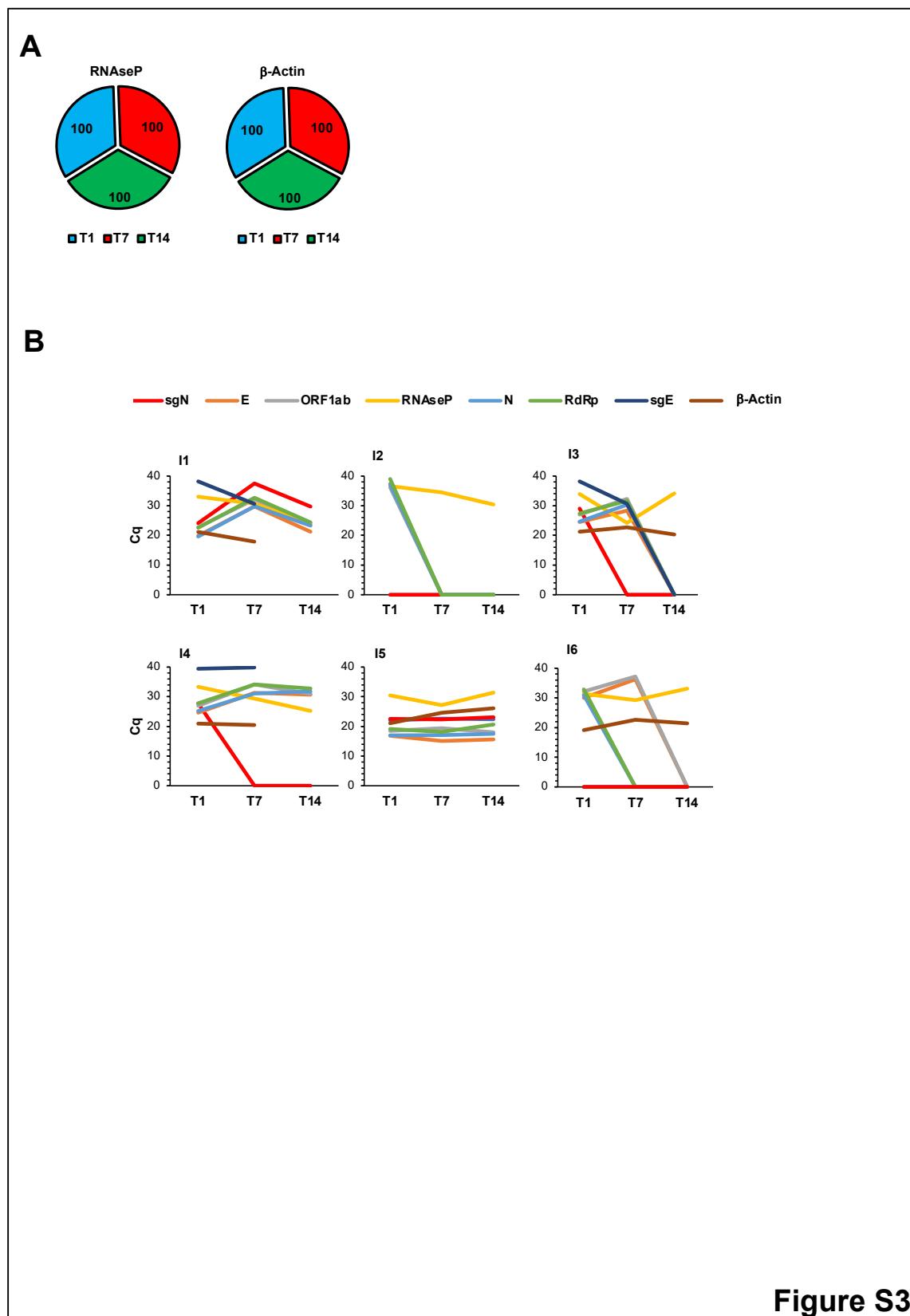


Figure S2

Supplementary Figure S2. Loss of detection of sgN precedes SARS-CoV-2 replication failure in home-isolated COVID19-affected patients. Related to Figures 2A-B. **(A)** Pie charts showing the proportions (%) of positivity of the oro/nasopharyngeal samples from home-isolated patients to human RNase P and β -actin genes at the different times (blue, first swab [n = 16]; red, second swab collected after 3 days [n = 16]; green, third swab collected after 7 days [n = 10]). **(B)** Line graphs showing changes over time of the Cq values related to viral sgN (red), sgE (brown), gene N (green), gene E (orange), gene ORF1ab (light gray), RdRp (light blue), and human RNase P (yellow) and β -actin (dark gray) genes in the oro/nasopharyngeal samples from 16 home-isolated patients.

Supplementary Figure S3.



Supplementary Figure S3. Loss of detection of sgN precedes SARS-CoV-2 replication failure in home-isolated COVID19-affected patients. Related to Figures 2C-D. **(A)** Pie chart showing the proportions (%) of positivity of the oro/nasopharyngeal samples from 6 hospitalized COVID19-positive patients to human RNase P gene at the different times (blue, first swab; red, second swab collected after 7 days; green, third swab collected after 14 days). **(B)** Line graphs showing the changes over time of the Cq values related to viral sgN (red), gene N (light blue), gene E (orange), gene ORF1ab (light gray), and RdRp (green), human RNase P gene (yellow), sgE (dark blue and orange) and β -Actin (brown) in the oro/nasopharyngeal samples from 6 hospitalized patients.

Supplementary Figure S4.

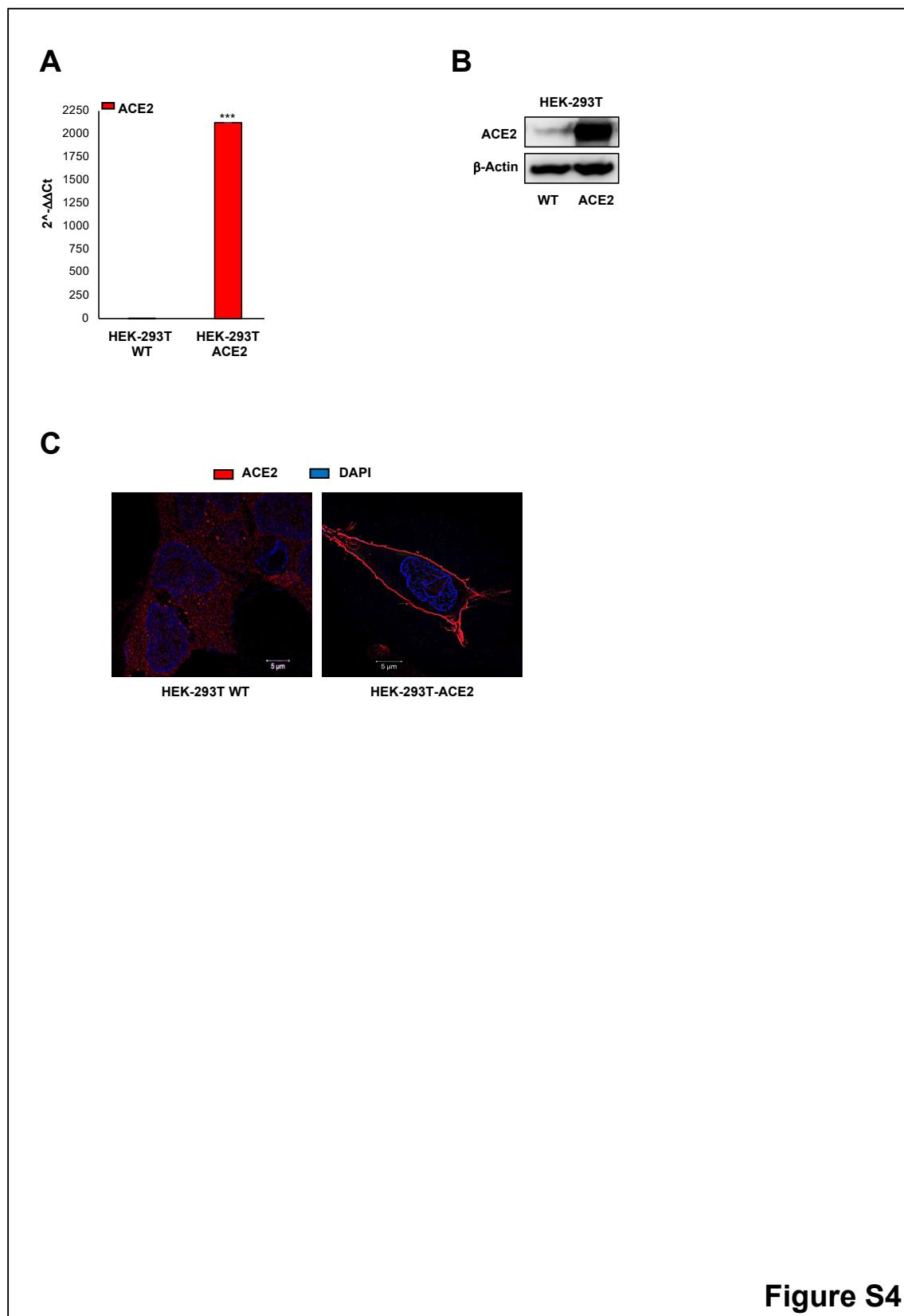


Figure S4

Supplementary Figure S4. HEK-293T overexpressing ACE2 cellular model. Related to Figures 3C-E, 4A-B. **(A)** Quantification of mRNA abundance relative to that of HEK-293T cells ($2^{-\Delta\Delta Ct}$) for the ACE2 gene. RT-PCR analysis with SYBR Green of RNA extracted from HEK-293T wild-type (WT) and HEK-293T—overexpressing ACE2 cells. Data are means \pm SD. ***P <0.001 (unpaired two-tailed Student's t test; n = 3 independent experiments per group). **(B)** Representative immunoblotting (using antibodies against the indicated proteins) of human HEK-293T cells and HEK-293T stable clones overexpressing ACE2. B-Actin was used as the loading control. **(C)** Immunofluorescence staining with an antibody against the human ACE2 (red) protein in human HEK-293T cells and HEK-293T stable clones overexpressing ACE2. DAPI was used for nuclei. The SIM image was acquired with Elyra 7 and processed with the Zeiss ZEN software (blue edition). Magnification, $\times 63$. Scale bars: 5 μ m.

Supplementary Figure S5.

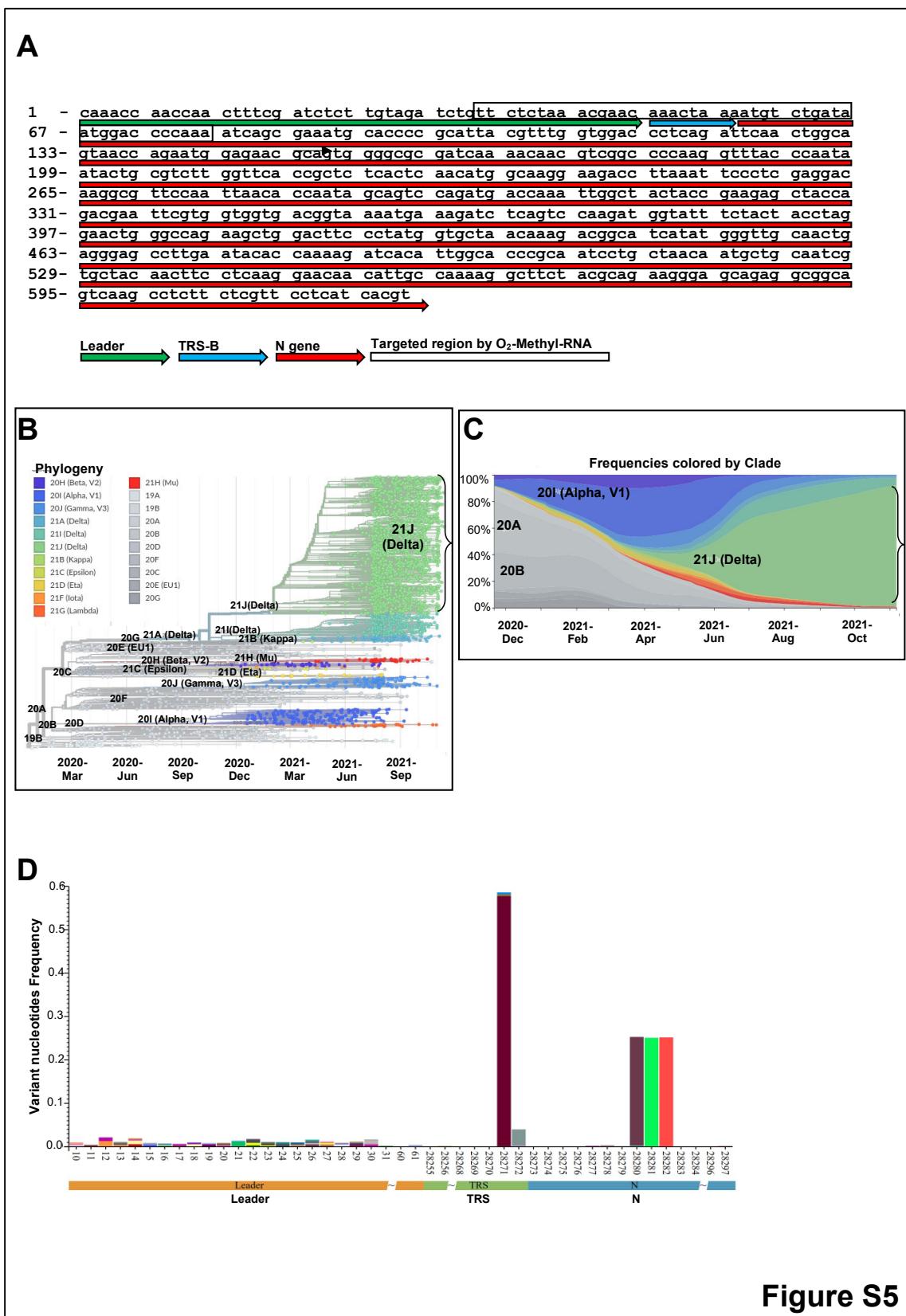


Figure S5

Supplementary Figure S5. 2'-O-methyl RNA against sgN sequence. Related to Figure 3A-B. **(A)** Sanger sequencing of the sgN region of SARS-CoV-2. Green arrow indicates the leader sequence, blue arrow indicates the TRS upstream gene N, and red arrow indicates gene N. The region recognized by 2'-O-methyl antisense RNA is highlighted in yellow. **(B, C)** Clade 21J is now the predominant form of Delta, with an estimated ~90% global frequency. **(D)** Frequencies of variant nucleotides calculated on a collection of about 350,000 sequences. Dataset downloaded from Global Initiative on Sharing Avian Influenza Data (GISAID) on November 15, 2021.

Supplementary Tables

Supplementary Table S1: Related to Figure 1. Cq values obtained for 50 oro/nasopharyngeal swabs from COVID19-positive patients according to the SARS-CoV-2 Viral3 kit and the Allplex 2019-nCoV assay.

Code	Cq value						
	SARS-CoV-2 Viral3 kit				Allplex 2019-nCoV assay		
	E gene	sgN transcript	ORF1ab gene	RNase P gene	E gene	N gene	RdRP/S gene
S8	16.36	20.37	15.83	33.28	13.84	15.58	14.16
S48	17.78	25.60	16.79	30.22	15.51	16.66	15.62
S4	15.15	18.56	14.54	30.56	12.08	13.38	15.65
S82	20.66	28.96	19.76	36.21	17.06	18.20	16.05
S37	19.38	24.17	18.49	30.75	13.30	15.67	16.64
S94	19.04	30.32	19.50	30.84	13.50	14.83	16.69
S16	16.31	22.26	16.20	31.43	13.79	15.73	16.74
S162	19.17	35.75	19.19	33.42	17.55	18.53	17.07
S33	18.88	23.82	17.82	30.80	17.34	18.56	17.18
S36	16.28	24.12	15.33	35.34	13.69	16.79	17.29
S35	17.26	24.02	16.62	31.80	16.60	16.48	17.33
S46	17.84	25.59	17.80	34.92	16.84	18.22	17.42
S43	19.37	24.63	18.52	31.58	16.88	18.78	17.46
S188	20.87	38.78	24.66	29.88	15.99	15.95	17.63
S6	16.54	19.28	16.00	31.48	16.58	17.68	17.67
S5	18.67	18.82	17.66	32.43	17.03	18.00	17.97
S11	15.80	20.82	16.31	31.15	15.78	15.92	18.34
S32	20.59	23.60	20.54	30.31	18.59	19.18	18.43
S23	18.47	22.73	17.53	32.01	16.74	16.91	18.53
S10	18.45	20.66	19.65	24.36	17.95	18.14	18.57
S27	20.58	23.31	19.50	31.98	19.04	19.77	18.75
S3	16.73	17.83	16.03	31.71	17.20	17.73	18.82
S25	18.71	23.09	19.06	34.34	18.68	19.87	18.88
S7	18.05	19.90	17.58	34.08	17.03	17.27	18.94
S62	21.45	26.65	23.51	38.09	19.23	20.55	19.09
S113	22.67	31.48	22.24	33.66	19.30	20.53	19.16
S59	22.20	26.50	21.13	34.97	19.56	21.16	19.28
S28	18.31	23.41	17.27	29.76	17.58	18.50	19.30
S29	18.75	23.56	19.09	34.11	17.90	17.60	19.68
S69	18.38	27.49	18.18	29.52	17.82	18.52	19.71
S42	19.15	24.57	19.19	32.09	19.07	20.46	20.07
S34	19.01	23.86	19.17	32.48	18.68	18.33	20.19
S100	22.07	30.67	21.77	33.91	20.54	21.73	20.37
S64	20.18	26.96	19.73	34.76	19.08	19.38	20.67
S9	18.78	20.63	18.21	31.34	18.63	18.67	20.70
S60	21.18	26.51	19.89	35.12	20.82	19.64	20.71
S70	21.18	27.65	21.16	38.10	18.84	19.21	20.94
S12	19.46	20.96	19.64	31.86	18.83	19.03	21.21

S61	19.59	26.54	19.72	33.18	19.43	19.54	21.25
S67	21.28	27.30	21.02	34.16	19.69	19.86	21.39
S73	26.40	27.75	26.50	35.30	24.31	24.95	23.14
S84	24.14	29.04	24.19	32.04	23.95	24.12	25.81
S120	28.05	31.87	28.04	36.70	28.22	28.68	26.32
S106	25.72	31.05	25.38	32.90	25.08	26.34	26.45
S163	30.14	35.75	29.64	36.23	28.30	27.76	26.56
S121	31.98	31.90	30.87	38.98	28.59	27.68	26.91
S156	29.34	35.06	29.15	35.59	28.79	28.89	27.15
S119	27.28	31.84	27.44	34.63	25.27	25.79	27.16
S96	28.64	30.49	28.91	27.29	26.11	25.69	28.21
S161	30.29	35.39	30.30	33.30	27.03	27.30	28.22

Supplementary Table S2. Related to Figure 1. Cq values obtained for 12 oro/nasopharyngeal swabs from negative COVID19-tested people according to the SARS-CoV-2 Viral3 kit and the Allplex 2019-nCoV assay.

Code	Cq values						
	SARS-CoV-2 Viral3 kit				Allplex 2019-nCoV assay		
	E gene	sgN transcript	ORF1ab gene	RNase P gene	E gene	N gene	RdRP/S gene
N1	N.A.	N.A.	N.A.	38.26	N.A.	N.A.	N.A.
N2	N.A.	N.A.	N.A.	34.59	N.A.	N.A.	N.A.
N3	N.A.	N.A.	N.A.	34.82	N.A.	N.A.	N.A.
N4	N.A.	N.A.	N.A.	31.12	N.A.	N.A.	N.A.
N6	N.A.	N.A.	N.A.	37.75	N.A.	N.A.	N.A.
N7	N.A.	N.A.	N.A.	25.58	N.A.	N.A.	N.A.
N8	N.A.	N.A.	N.A.	28.54	N.A.	N.A.	N.A.
N9	N.A.	N.A.	N.A.	23.17	N.A.	N.A.	N.A.
N10	N.A.	N.A.	N.A.	26.42	N.A.	N.A.	N.A.
N11	N.A.	N.A.	N.A.	30.46	N.A.	N.A.	N.A.
N12	N.A.	N.A.	N.A.	26.49	N.A.	N.A.	N.A.

N.A., not amplified (i.e., Cq value >40).

Supplementary Table S3. Related to Figure 1A-B. Cq values obtained from the 315 oro/nasopharyngeal swabs from COVID19-positive patients with the SARS-CoV-2 Viral3 kit.

Code	Cq value				Code	Cq value			
	E	sgN	ORF1ab	RNase P		E	sgN	ORF1ab	RNAse P
S1	16.18	15.12	16.41	28.45	S159	26.61	35.31	26.44	27.75
S2	20.43	17.02	21.45	29.23	S160	26.61	35.31	26.44	27.75
S3	16.73	17.83	16.03	31.71	S161	30.29	35.39	30.30	33.30
S4	15.15	18.56	14.54	30.56	S162	19.17	35.75	19.19	33.42
S5	18.67	18.82	17.66	32.43	S163	30.14	35.75	29.64	36.23
S6	16.54	19.28	16.00	31.48	S164	28.31	36.57	29.10	26.02
S7	18.05	19.90	17.58	34.08	S165	28.31	36.57	29.10	26.02
S8	16.36	20.37	15.83	33.28	S166	29.16	36.59	28.94	28.64
S9	18.78	20.63	18.21	31.34	S167	30.27	36.62	30.73	29.67
S10	18.45	20.66	19.65	24.36	S168	14.74	36.77	14.09	22.18
S11	15.80	20.82	16.31	31.15	S169	14.74	36.77	14.09	22.18
S12	19.46	20.96	19.64	31.86	S170	25.85	36.94	25.22	28.00
S13	15.12	21.79	17.46	26.59	S171	29.05	36.98	29.18	26.40
S14	15.12	21.79	17.46	26.59	S172	26.98	37.00	26.67	32.00
S15	15.16	22.05	14.65	27.00	S173	26.66	37.19	25.42	23.01
S16	16.31	22.26	16.20	31.43	S174	26.66	37.19	25.42	23.01
S17	17.70	22.34	18.73	26.82	S175	29.90	37.24	30.46	25.64
S18	17.70	22.34	18.73	26.82	S176	29.90	37.24	30.46	25.64
S19	17.35	22.41	17.52	24.75	S177	27.62	37.42	27.43	32.00
S20	17.35	22.41	17.52	24.75	S178	30.77	37.56	31.18	31.61
S21	20.04	22.61	20.32	26.64	S179	30.77	37.56	31.18	31.61
S22	20.04	22.61	20.32	26.64	S180	27.13	37.60	27.27	32.08
S23	18.47	22.73	17.53	32.01	S181	33.59	37.72	34.24	30.88
S24	23.13	23.07	23.73	28.47	S182	30.40	37.80	31.35	31.67
S25	18.71	23.09	19.06	34.34	S183	25.01	37.89	25.46	29.65
S26	17.70	21.43	17.94	29.25	S184	25.01	37.89	25.46	29.65
S27	20.58	23.31	19.50	31.98	S185	28.16	37.98	28.84	31.13
S28	18.31	23.41	17.27	29.76	S186	28.16	37.98	28.84	31.13
S29	18.75	23.56	19.09	34.11	S187	22.38	38.25	23.04	26.27
S30	18.87	23.59	19.13	26.55	S188	20.87	38.78	24.66	29.88
S31	18.87	23.59	19.13	26.55	S189	28.56	38.85	28.01	28.93
S32	20.59	23.60	20.54	30.31	S190	26.06	38.88	25.22	30.53
S33	18.88	23.82	17.82	30.80	S191	30.17	38.90	29.02	32.24
S34	19.01	23.86	19.17	32.48	S192	32.64	38.99	32.48	34.09
S35	17.26	24.02	16.62	31.80	S193	30.14	39.02	30.21	33.30
S36	16.28	24.12	15.33	35.34	S194	30.34	39.12	30.15	32.11
S37	19.38	24.17	18.49	30.75	S195	27.54	39.19	26.83	33.10
S38	20.05	24.22	20.97	30.60	S196	24.32	N.A.	23.17	25.21
S39	20.05	24.22	20.97	30.60	S197	26.50	N.A.	25.73	31.21
S40	21.42	24.40	22.02	26.14	S198	26.52	N.A.	26.52	26.43
S41	21.42	24.40	22.02	26.14	S199	26.63	N.A.	25.84	28.65
S42	19.15	24.57	19.19	32.09	S200	26.88	N.A.	27.61	33.86
S43	19.37	24.63	18.52	31.58	S201	26.94	N.A.	26.39	30.14
S44	20.51	25.03	20.63	28.18	S202	27.02	N.A.	25.57	25.00

S45	20.52	25.43	21.19	30.08	S203	27.06	N.A.	26.25	31.53
S46	17.84	25.59	17.80	34.92	S204	27.10	N.A.	26.11	27.56
S47	19.26	23.94	19.15	30.60	S205	27.15	N.A.	26.35	30.88
S48	17.78	25.60	16.79	30.22	S206	28.01	N.A.	28.08	21.13
S49	20.33	25.71	21.10	29.35	S207	28.12	N.A.	26.85	22.56
S50	19.37	25.78	19.73	25.70	S208	28.15	N.A.	29.20	31.10
S51	17.01	25.88	17.19	22.79	S209	28.54	N.A.	27.03	23.46
S52	17.01	25.88	17.19	22.79	S210	29.44	N.A.	28.43	21.68
S53	25.46	25.98	25.57	30.39	S211	29.47	N.A.	29.67	32.39
S54	19.61	26.00	20.50	30.43	S212	29.48	N.A.	29.80	32.75
S55	20.33	26.11	21.56	28.38	S213	29.66	N.A.	29.36	31.44
S56	20.15	26.15	20.81	28.35	S214	30.18	N.A.	30.60	28.63
S57	17.47	26.39	17.99	27.70	S215	30.35	N.A.	31.17	34.03
S58	17.47	26.39	17.99	27.70	S216	30.73	N.A.	29.51	25.68
S59	22.20	26.50	21.13	34.97	S217	30.77	N.A.	31.03	32.09
S60	21.18	26.51	19.89	35.12	S218	30.86	N.A.	29.52	31.64
S61	19.59	26.54	19.72	33.18	S219	31.10	N.A.	32.20	29.02
S62	21.45	26.65	23.51	38.09	S220	31.22	N.A.	31.75	32.61
S63	20.04	26.70	20.05	31.00	S221	31.29	N.A.	31.42	31.98
S64	20.18	26.96	19.73	34.76	S222	31.39	N.A.	32.22	24.57
S65	21.38	27.24	22.25	26.96	S223	31.39	N.A.	31.47	32.37
S66	21.38	27.24	22.25	26.96	S224	31.45	N.A.	31.36	32.47
S67	21.28	27.30	21.02	34.16	S225	31.47	N.A.	31.92	31.60
S68	22.07	27.40	21.94	31.11	S226	31.51	N.A.	32.14	33.78
S69	18.38	27.49	18.18	29.52	S227	31.63	N.A.	31.82	34.04
S70	21.18	27.65	21.16	38.10	S228	31.69	N.A.	31.58	32.29
S71	21.60	27.70	21.03	30.09	S229	31.71	N.A.	31.85	30.55
S72	18.59	27.74	25.70	27.47	S230	31.95	N.A.	30.63	23.50
S73	26.40	27.75	26.50	35.30	S231	32.00	N.A.	32.02	26.08
S74	20.46	27.88	20.10	31.23	S232	32.15	N.A.	31.31	23.07
S75	20.67	28.11	22.61	29.13	S233	32.16	N.A.	33.77	26.18
S76	20.15	28.13	20.94	28.82	S234	32.20	N.A.	32.54	35.03
S77	20.15	28.13	20.94	28.82	S235	32.22	N.A.	32.35	30.50
S78	19.28	28.20	19.32	27.46	S236	32.40	N.A.	33.78	34.06
S79	24.13	28.64	24.52	29.42	S237	32.40	N.A.	33.23	33.86
S80	18.59	28.75	17.64	25.18	S238	32.53	N.A.	32.19	27.44
S81	18.59	28.75	17.64	25.18	S239	32.69	N.A.	32.55	23.08
S82	20.66	28.96	19.76	36.21	S240	32.75	N.A.	32.02	31.48
S83	16.36	29.02	16.64	19.82	S241	32.75	N.A.	32.76	33.57
S84	24.14	29.04	24.19	32.04	S242	32.89	N.A.	33.02	27.95
S85	24.92	29.23	24.69	31.00	S243	32.98	N.A.	33.18	31.03
S86	18.45	29.29	26.88	28.45	S244	33.05	N.A.	31.81	27.02
S87	18.75	29.52	19.69	20.76	S245	33.10	N.A.	32.11	31.61
S88	24.25	29.53	24.07	31.00	S246	33.12	N.A.	33.02	28.33
S89	23.79	29.66	23.34	31.90	S247	33.13	N.A.	33.20	31.20
S90	24.32	29.82	24.83	27.84	S248	33.15	N.A.	33.43	30.83
S91	18.52	30.18	17.72	25.35	S249	33.15	N.A.	32.97	32.37
S92	18.52	30.18	17.72	25.35	S250	33.23	N.A.	32.75	32.23
S93	22.44	30.24	22.75	30.25	S251	33.25	N.A.	32.22	26.29
S94	19.04	30.32	19.50	30.84	S252	33.30	N.A.	33.04	33.50

S95	16.89	30.42	18.91	19.71	S253	33.34	N.A.	34.05	29.96
S96	28.64	30.49	28.91	27.29	S254	33.38	N.A.	33.56	30.59
S97	24.37	30.49	24.43	32.00	S255	33.44	N.A.	33.68	25.23
S98	23.24	30.49	22.81	32.08	S256	33.49	N.A.	33.07	27.39
S99	22.06	30.51	30.21	29.12	S257	33.67	N.A.	33.03	34.98
S100	22.07	30.67	21.77	33.91	S258	33.70	N.A.	32.72	21.89
S101	25.61	30.69	25.58	33.29	S259	33.74	N.A.	33.96	31.10
S102	21.08	30.76	20.53	29.81	S260	33.84	N.A.	34.09	26.58
S103	23.30	30.76	23.85	30.39	S261	33.85	N.A.	32.48	29.74
S104	21.36	30.84	22.62	22.84	S262	33.95	N.A.	33.95	26.49
S105	22.56	31.05	23.08	31.70	S263	34.09	N.A.	33.69	34.35
S106	25.72	31.05	25.38	32.90	S264	34.10	N.A.	33.74	28.07
S107	19.69	31.06	29.08	28.42	S265	34.10	N.A.	33.35	30.66
S108	28.16	31.07	28.53	32.00	S266	34.14	N.A.	35.26	34.66
S109	22.14	31.11	27.65	29.21	S267	34.19	N.A.	35.11	28.51
S110	23.80	31.27	23.59	30.10	S268	34.23	N.A.	34.87	30.28
S111	18.95	31.29	19.45	23.45	S269	34.25	N.A.	34.61	22.51
S112	26.14	31.42	25.98	28.36	S270	34.25	N.A.	32.92	31.49
S113	22.67	31.48	22.24	33.66	S271	34.25	N.A.	35.98	28.15
S114	21.48	31.58	22.66	25.70	S272	34.26	N.A.	35.01	30.14
S115	22.10	31.65	21.05	21.73	S273	34.33	N.A.	33.62	26.48
S116	22.10	31.65	21.05	21.73	S274	34.36	N.A.	35.03	32.93
S117	24.28	31.68	25.58	31.35	S275	34.37	N.A.	34.51	20.43
S118	25.25	31.70	25.24	32.01	S276	34.48	N.A.	35.23	32.24
S119	27.28	31.84	27.44	34.63	S277	34.50	N.A.	34.41	31.16
S120	28.05	31.87	28.04	36.70	S278	34.50	N.A.	34.60	31.16
S121	31.98	31.90	30.87	38.98	S279	34.59	N.A.	34.40	33.33
S122	25.74	31.98	26.20	30.02	S280	34.68	N.A.	33.89	30.21
S123	24.21	32.09	23.83	30.06	S281	34.73	N.A.	34.49	34.39
S124	23.43	32.25	23.94	28.28	S282	34.73	N.A.	35.13	33.30
S125	26.33	32.25	26.27	31.44	S283	34.90	N.A.	35.58	31.01
S126	20.19	32.36	20.10	27.42	S284	35.06	N.A.	34.38	27.45
S127	25.41	32.40	24.54	29.38	S285	35.12	N.A.	33.81	21.71
S128	21.69	32.46	24.69	23.94	S286	35.12	N.A.	37.19	30.61
S129	26.89	32.53	26.45	29.14	S287	35.16	N.A.	34.67	7.06
S130	20.78	32.58	19.59	25.15	S288	35.18	N.A.	35.74	25.19
S131	20.78	32.58	19.59	25.15	S289	35.20	N.A.	36.04	30.15
S132	24.49	32.68	26.02	28.34	S290	35.34	N.A.	36.49	31.28
S133	26.83	32.83	31.92	33.50	S291	35.40	N.A.	35.57	26.87
S134	18.68	32.85	18.97	26.55	S292	35.47	N.A.	35.91	35.66
S135	29.20	32.97	29.47	38.03	S293	35.76	N.A.	36.34	32.81
S136	27.45	33.04	29.30	29.65	S294	35.83	N.A.	37.33	28.48
S137	27.57	33.06	27.97	30.07	S295	35.94	N.A.	38.10	30.48
S138	25.67	33.32	26.17	31.17	S296	36.00	N.A.	36.58	32.74
S139	25.67	33.32	26.17	31.17	S297	36.08	N.A.	34.14	32.90
S140	23.63	33.35	23.65	23.13	S298	36.13	N.A.	37.50	29.16
S141	23.63	33.35	23.65	23.13	S299	36.16	N.A.	36.35	31.19
S142	24.67	33.44	31.25	33.90	S300	36.18	N.A.	36.31	30.95
S143	24.86	33.45	23.99	30.00	S301	36.21	N.A.	36.70	31.10
S144	27.30	33.53	27.19	28.84	S302	36.28	N.A.	35.74	34.68

S145	20.64	33.71	21.34	24.45	S303	36.39	N.A.	37.25	26.97
S146	23.56	33.72	23.53	30.08	S304	36.71	N.A.	38.21	26.99
S147	20.34	33.77	19.20	23.64	S305	36.74	N.A.	38.25	35.29
S148	20.34	33.77	19.20	23.64	S306	36.98	N.A.	36.69	28.51
S149	24.95	33.92	25.60	33.28	S307	37.13	N.A.	36.05	31.11
S150	27.78	33.96	28.00	31.88	S308	37.19	N.A.	38.09	31.35
S151	27.72	34.07	27.81	30.14	S309	37.61	N.A.	36.69	28.06
S152	28.87	34.18	29.79	33.89	S310	38.07	N.A.	36.32	30.28
S153	22.87	34.24	21.34	21.81	S311	38.30	N.A.	40.20	25.99
S154	22.87	34.24	21.34	21.81	S312	39.79	N.A.	38.81	35.17
S155	22.07	34.58	21.88	31.90	S313	39.79	N.A.	37.27	29.62
S156	29.34	35.06	29.15	35.59	S314	40.23	N.A.	40.40	30.22
S157	19.16	35.14	19.35	26.84	S315	40.26	N.A.	40.08	30.52
S158	19.16	35.14	19.35	26.84					

N.A., not amplified (i.e., Cq >40)

Supplementary Table S4. Related to Figure 1D,F,H. Cq values for sgE and β-actin obtained from the single cohort of 122 oro/nasopharyngeal swabs from COVID19-positive patients analyzed using Taqman qPCR.

Code	Cq value										
	sgE	β-actin									
S44	27.37	26.50	S312	39.40	26.95	S261	N.A.	24.48	S274	N.A.	25.44
S45	27.50	27.75	S191	39.49	26.34	S239	N.A.	26.2	S267	N.A.	27.54
S49	26.92	25.39	S200	39.65	28.35	S250	N.A.	30.57	S276	N.A.	28.45
S54	26.19	27.27	S199	39.74	29.76	S241	N.A.	30.78	S266	N.A.	33.50
S55	27.36	25.92	S219	39.88	28.03	S270	N.A.	28.09	S291	N.A.	23.98
S56	26.80	25.39	S243	38.04	27.35	S249	N.A.	29.90	S283	N.A.	28.03
S71	30.75	28.07	S256	38.25	32.94	S242	N.A.	29.51	S302	N.A.	27.18
S74	40.4	30.28	S212	38.57	29.13	S246	N.A.	27.11	S288	N.A.	25.24
S98	31.27	28.53	S215	38.81	29.84	S257	N.A.	24.47	S292	N.A.	33.38
S105	37.98	28.06	S213	39.17	29.19	S252	N.A.	26.87	S271	N.A.	25.56
S112	35.15	27.05	S280	39.45	27.76	S247	N.A.	29.69	S289	N.A.	26.91
S127	36.95	30.51	S227	38.73	30.58	S237	N.A.	30.55	S307	N.A.	28.49
S137	37.45	27.42	S234	38.34	31.15	S265	N.A.	29.04	S300	N.A.	27.74
S144	40.10	28.67	S203	N.A.	28.14	S248	N.A.	30.38	S310	N.A.	24.03
S149	33.92	27.80	S201	N.A.	29.33	S254	N.A.	27.94	S293	N.A.	30.41
S150	37.49	28.96	S198	N.A.	27.84	S273	N.A.	25.37	S299	N.A.	27.48
S151	34.30	26.84	S218	N.A.	27.33	S263	N.A.	31.51	S290	N.A.	22.25
S166	38.85	26.63	S217	N.A.	30.21	S264	N.A.	27.97	S306	N.A.	30.13
S167	40.10	26.15	S224	N.A.	30.52	S262	N.A.	24.45	S301	N.A.	28.23
S180	41.10	30.08	S221	N.A.	27.62	S259	N.A.	25.81	S286	N.A.	28.38
S190	38.03	26.79	S223	N.A.	29.47	S253	N.A.	26.29	S303	N.A.	27.48
S238	28.94	25.97	S228	N.A.	29.21	S260	N.A.	25.81	S294	N.A.	22.79
S206	31.03	28.27	S220	N.A.	27.22	S297	N.A.	30.51	S298	N.A.	27.30
S275	34.37	26.10	S244	N.A.	29.69	S284	N.A.	29.72	S308	N.A.	28.20
S282	36.72	28.70	S229	N.A.	25.35	S279	N.A.	27.47	S295	N.A.	27.27
S208	37.82	27.04	S225	N.A.	30.02	S277	N.A.	30.19	S304	N.A.	25.69
S205	38.18	28.11	S231	N.A.	30.60	S281	N.A.	33.87	S305	N.A.	30.99
S211	38.47	27.09	S240	N.A.	29.44	S278	N.A.	27.52	S315	N.A.	29.07
S197	38.87	28.86	S245	N.A.	28.46	S287	N.A.	25.94	S311	N.A.	31.67
S236	39.40	30.65	S226	N.A.	30.72	S268	N.A.	26.54			
S235	N.A.	29.60	S272	N.A.	29.32	S314	N.A.	25.23			

N.A., not amplified (i.e., Cq >40)

Supplementary Table S5. Related to Figure 2A-B. Cq values obtained from a cohort of 16 oro/nasopharyngeal swabs from home-isolated COVID19-infected patients analyzed according to scheduled times (i.e., 3-day intervals from the first swab) with the SARS-CoV-2 Viral3 kits (sgN, E gene, ORF1ab gene) and the Allplex 2019-nCoV assay (E, N, RdRp/S genes) and by Taqman qPCR (sgE, β-actin genes).

Marker	Day	Cq value															
		#1	#2	#3	#4	#5	#6	#7	#8	#9	#10	#11	#12	#19	#20	#21	#22
SARS-CoV-2 Viral3 kit																	
sgN transcript	1	29.87	33.08	36.46	37.16	29.21	21.38	20.92	16.7	30.58	29.67	31.07	17.38	26.22	24.26	27.07	25.41
	3	N.A.	N.A.	N.A.	N.A.	N.A.	26.59	36.35	32.29	37.87	23.25	29.71	32.58	N.A.	N.A.	N.A.	N.A.
	7	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	24.95	-	-	-	-	-	-	-
E gene	1	24.22	28.52	29.34	34.5	23.89	21.38	20.15	13.3	25.13	16.44	23.96	12.75	21.23	20.57	20.77	18.25
	3	30.43	36.28	30.76	28.04	31.49	18.16	29.03	24.76	29.5	15.54	23.32	17.48	33.87	35.23	38.28	37.23
	7	N.A.	N.A.	N.A.	N.A.	N.A.	33.78	N.A.	N.A.	17.83	-	-	-	-	-	-	-
Orflab gene	1	26.77	31.55	31.88	35.04	24.84	23.62	23.18	15.05	30.85	18.47	27.18	11.34	22.51	22.39	21.15	19.68
	3	34.84	34.94	35.95	30.98	33.37	20.98	35.83	25.87	32.32	18.37	25.99	25.67	37.03	36.38	39.65	38.64
	7	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	37.91	18.52	-	-	-	-	-	-	-
RNase P gene	1	32.23	28.27	33.49	32.14	30.91	27.89	31.21	28.34	31	30.13	31.11	31.42	29.11	31.04	27.53	29.93
	3	31.2	32.07	29.46	34.11	28.87	31.07	26.31	30.24	32.87	30.02	31.86	31.49	29.56	31.39	31.37	32.42
	7	31.7	32.4	30.23	30.01	36.56	32.25	34	32.1	32.87	27.87	-	-	-	-	-	-
Allplex 2019-nCoV assay																	
RdRp/S gene	1	26.93	32.94	26.55	34.94	25.45	25.64	24.46	23.78	31.55	27.03	29.61	21.26	22.47	24.71	25.03	20.05
	3	31.34	27.15	34.6	38.74	33.12	20.38	31.23	29.84	34.64	17.41	27.64	30.67	37.51	31.98	37.86	37.49
	7	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	N.A.	17.66	-	-	-	-	-	-	-
N gene	1	23.96	28.35	21.91	35.81	27.44	21.60	19.90	18.94	26.92	19.03	25.38	21.09	24.25	21.85	24.6	19.02
	3	31.5	37.01	30.55	35.56	34.42	20.59	30.17	27.42	31.05	17.6	25.14	29.98	32.97	30.09	38.71	35.45
	7	N.A.	N.A.	N.A.	N.A.	37.23	36	N.A.	N.A.	18.21	-	-	-	-	-	-	-
E gene	1	24.55	29.38	21.74	34.37	24.98	22.30	21.19	19.04	26.62	20.71	24.79	20.14	22.06	21.01	23.37	17.54
	3	32.29	38.74	31.03	35.52	31.26	20.54	31.23	25.6	31.39	17.3	24.78	32.35	33.06	30.18	39.22	35.86
	7	N.A.	N.A.	N.A.	N.A.	N.A.	36.05	N.A.	N.A.	17.94	-	-	-	-	-	-	-
Taqman qPCR																	
sgE transcript	1	29.84	33.08	26.11	-	29.55	27.78	24.11	33.17	31.63	-	-	-	-	-	-	-
	3	36.42	38.97	35.01	-	38.4	29.18	35.68	37.35	35.64	-	-	-	-	-	-	-
	7	N.A.	N.A.	N.A.	-	N.A.	N.A.	38.81	N.A.	N.A.	-	-	-	-	-	-	-
β-actin gene	1	26.42	22.96	23.42	-	28.45	22.42	25.68	23.52	27.42	-	-	-	-	-	-	-
	3	23.63	26.83	27.39	-	26.53	24.21	24.39	26.11	26.18	-	-	-	-	-	-	-
	7	31.2	26.58	24.72	-	31.23	24.73	25.99	30.44	25.97	-	-	-	-	-	-	-

N.A., not amplified (i.e., Cq >40)

Supplementary Table S6. Related to Figure 2C-D. Cq values obtained from a cohort of 6 oro/nasopharyngeal swabs from hospitalized COVID19-positive patients analyzed according to scheduled times (i.e., 7-day intervals from the first swab) with the SARS-CoV-2 Viral3 kits (sgN, E gene, ORF1ab gene) and the Allplex 2019-nCoV assay (E, N, RdRp/S genes) and by Taqman qPCR (sgE, β-actin genes).

Marker	Days	Cq value					
		I1	I2	I3	I4	I5	I6
SARS-CoV-2 Viral3 kits							
sgN transcript	1	24.13	N.A.	29.01	27.76	22.06	N.A.
	7	37.47	N.A.	N.A.	N.A.	22.62	N.A.
	14	29.63	N.A.	N.A.	N.A.	22.44	N.A.
E gene	1	19.85	37.36	24.56	24.66	16.86	30.04
	7	29.76	N.A.	28.32	31.32	15.09	36.27
	14	21.24	N.A.	N.A.	30.67	15.64	N.A.
ORF1ab gene	1	22.79	36.2	26.95	26.98	18.47	32.16
	7	32.11	N.A.	32.31	34.18	19.37	37.2
	14	23.49	N.A.	N.A.	31.03	18.01	N.A.
RNaseP	1	33.01	36.6	33.93	33.31	30.44	31.34
	7	30.63	34.5	24.21	29.39	27.2	29.24
	14	24.05	30.37	34.12	25.22	31.42	33.14
Allplex 2019-nCoV assay							
E gene	1	20.55	38.46	25.44	25.85	17.26	30.44
	7	30.27	N.A.	29.34	32.1	16.26	37.76
	14	21.96	N.A.	N.A.	30.79	18	N.A.
N gene	1	19.57	37.1	24.61	25.21	17.01	30.92
	7	29.8	N.A.	30.31	31.17	17.11	N.A.
	14	23.25	N.A.	N.A.	31.77	17.52	N.A.
RdRp/S gene	1	22.54	39	27.43	27.8	19.12	32.88
	7	32.69	N.A.	31.64	34.16	18.13	N.A.
	14	24.32	N.A.	N.A.	32.71	20.67	N.A.
Taqman qPCR							
sgE transcript	1	38.15	-	39.13	39.45	22.58	39.55
	7	30.59	-	39.88	39.86	22.34	N.A.
	14	-	-	N.A.	-	23.09	N.A.
β-actin gene	1	21.15	-	21.22	20.93	21.07	19.16
	7	17.87	-	22.68	20.49	24.61	22.64
	14	-	-	20.28	-	26.08	21.41

N.A., not amplified (i.e., Cq values >40)