

REFERENCES

- Amarasinghe, S. L., Su, S., Dong, X., Zappia, L., Ritchie, M. E., & Gouil, Q. (2020). Opportunities and challenges in long-read sequencing data analysis. *Genome Biology*, 21(1). <https://doi.org/10.1186/s13059-020-1935-5>
- Artic Network. (2021, December 1). *SARS-CoV-2 V4.1 update for Omicron variant*. ARTIC Real-Time Genomic Surveillance. <https://community.artic.network/t/sars-cov-2-v4-1-update-for-omicron-variant/342>
- Banerjee, A., Kulcsar, K., Misra, V., Frieman, M., & Mossman, K. (2019). Bats and Coronaviruses. *Viruses*, 11(1). <https://doi.org/10.3390/v11010041>
- Bio-Rad. (n.d.). *PCR Troubleshooting*. Bio-Rad Laboratories. <https://www.bio-rad.com/en-id/applications-technologies/pcr-troubleshooting?ID=LUSO3HC4S#:~:text=Annealing%20temperature%20was%20too%20high>
- Charre, C., Ginevra, C., Sabatier, M., Regue, H., Destras, G., Brun, S., Burfin, G., Scholtes, C., Morfin, F., Valette, M., Lina, B., Bal, A., & Josset, L. (2020). Evaluation of NGS-based approaches for SARS-CoV-2 whole genome characterisation. *Virus Evolution*, 6(2). <https://doi.org/10.1093/ve/veaa075>
- Chen, Z., Erickson, D. L., & Meng, J. (2021). Polishing the Oxford Nanopore long-read assemblies of bacterial pathogens with Illumina short reads to improve genomic analyses. *Genomics*, 113(3), 1366–1377. <https://doi.org/10.1016/j.ygeno.2021.03.018>
- Constantinides, B., Webster, H., Gentry, J., Bastable, J., Dunn, L., Oakley, S., Swann, J., Sanderson, N., Fowler, P. W., Ma, G., Rodger, G., Barrett, L., Jeffery, K., Peto, T. E., Stoesser, N., Street, T., & Crook, D. W. (2022). *Rapid turnaround multiplex sequencing of SARS-CoV-2: comparing tiling amplicon protocol performance*. <https://doi.org/10.1101/2021.12.28.21268461>
- Cruz-González, A., Muñoz-Velasco, I., Cottom-Salas, W., Becerra, A., Campillo-Balderas, J. A., Hernández-Morales, R., Vázquez-Salazar, A., Jácome, R., & Lazcano, A. (2021). Structural analysis of viral ExoN domains reveals polyphyletic hijacking events. *PLOS ONE*, 16(3), e0246981. <https://doi.org/10.1371/journal.pone.0246981>
- CSSE John Hopkins. (2022). *COVID-19 Data Repository by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University*. GitHub. <https://github.com/CSSEGISandData/COVID-19>
- Cucinotta, D., & Vanelli, M. (2020). WHO Declares COVID-19 a Pandemic. *Acta Bio-Medica: Atenei Parmensis*, 91(1), 157–160. <https://doi.org/10.23750/abm.v91i1.9397>
- Davis, J. J., Long, S. W., Christensen, P. A., Olsen, R. J., Olson, R., Shukla, M., Subedi, S., Stevens, R., & Musser, J. M. (2021). Analysis of the ARTIC Version 3 and Version 4 SARS-CoV-2 Primers and Their Impact on the Detection of the G142D Amino Acid Substitution in the Spike Protein. *Microbiology Spectrum*, 9(3). <https://doi.org/10.1128/spectrum.01803-21>
- Durmaz, B., Abdulmajed, O., & Durmaz, R. (2020). Mutations Observed in the SARS-CoV-2 Spike Glycoprotein and Their Effects in the Interaction of Virus with ACE-2 Receptor. *Medeniyet Medical Journal*. <https://doi.org/10.5222/mmj.2020.98048>
- EPI2ME Labs. (2022). *ARTIC SARS-CoV-2 Workflow*. GitHub. <https://github.com/epi2me-labs/wf-artic>
- Freed, N. E., Vlková, M., Faisal, M. B., & Silander, O. K. (2020). Rapid and Inexpensive Whole-Genome Sequencing of SARS-CoV-2 using 1200 bp Tiled Amplicons and Oxford Nanopore Rapid Barcoding. *Biology Methods and Protocols*. <https://doi.org/10.1093/biomethods/bpaa014>
- Freed, N., & Silander, O. (2021). SARS-CoV2 genome sequencing protocol (1200bp amplicon “midnight” primer set, using Nanopore Rapid kit) v2. *Protocols.io*. <https://doi.org/10.17504/protocols.io.bwypfvn>
- Gomes, A., & Korf, B. (2018). Genetic Testing Techniques. *Pediatric Cancer Genetics*, 47–64. <https://doi.org/10.1016/b978-0-323-48555-5.00005-3>
- González-Recio, O., Gutiérrez-Rivas, M., Peiró-Pastor, R., Aguilera-Sepúlveda, P., Cano-Gómez, C.,

- Jiménez-Clavero, M. Á., & Fernández-Pinero, J. (2021). Sequencing of SARS-CoV-2 genome using different nanopore chemistries. *Applied Microbiology and Biotechnology*, *105*(8), 3225–3234. <https://doi.org/10.1007/s00253-021-11250-w>
- Gorbacheva, T., Quispe-Tintaya, W., Popov, V. N., Vijg, J., & Maslov, A. Y. (2015). Improved transposon-based library preparation for the Ion Torrent platform. *BioTechniques*, *58*(4). <https://doi.org/10.2144/000114277>
- Hawkins, G. A. (2017). Analysis of Human Genetic Variations Using DNA Sequencing. *Basic Science Methods for Clinical Researchers*, 77–98. <https://doi.org/10.1016/b978-0-12-803077-6.00005-9>
- Itokawa, K., Sekizuka, T., Hashino, M., Tanaka, R., & Kuroda, M. (2020). Disentangling primer interactions improves SARS-CoV-2 genome sequencing by multiplex tiling PCR. *PLOS ONE*, *15*(9), e0239403. <https://doi.org/10.1371/journal.pone.0239403>
- Khailany, R. A., Safdar, M., & Ozaslan, M. (2020). Genomic characterization of a novel SARS-CoV-2. *Gene Reports*, *19*, 100682. <https://doi.org/10.1016/j.genrep.2020.100682>
- Kia, A., Gloeckner, C., Osothprarop, T., Gormley, N., Bomati, E., Stephenson, M., Goryshin, I., & He, M. M. (2017). Improved genome sequencing using an engineered transposase. *BMC Biotechnology*, *17*(1). <https://doi.org/10.1186/s12896-016-0326-1>
- Ku, C.-S., & Roukos, D. H. (2013). From next-generation sequencing to nanopore sequencing technology: paving the way to personalized genomic medicine. *Expert Review of Medical Devices*, *10*(1), 1–6. <https://doi.org/10.1586/erd.12.63>
- Lambisia, A. W., Mohammed, K. S., Makori, T. O., Ndwiga, L., Mburu, M. W., Morobe, J. M., Moraa, E. O., Musyoki, J., Murunga, N., Mwangi, J. N., Nokes, D. J., Agoti, C. N., Ochola-Oyier, L. I., & Githinji, G. (2022). Optimization of the SARS-CoV-2 ARTIC Network V4 Primers and Whole Genome Sequencing Protocol. *Frontiers in Medicine*, *9*. <https://doi.org/10.3389/fmed.2022.836728>
- Lee, H., Gurtowski, J., Yoo, S., Nattestad, M., Marcus, S., Goodwin, S., Richard McCombie, W., & Schatz, M. C. (2016). Third-generation sequencing and the future of genomics. *BioRxiv*. <https://doi.org/10.1101/048603>
- Ma, W., Yang, J., Fu, H., Su, C., Yu, C., Wang, Q., de Vasconcelos, A. T. R., Bazykin, G. A., Bao, Y., & Li, M. (2022). Genomic perspectives on the emerging SARS-CoV-2 omicron variant. *Genomics, Proteomics & Bioinformatics*. <https://doi.org/10.1016/j.gpb.2022.01.001>
- Mahase, E. (2020). Covid-19: WHO declares pandemic because of “alarming levels” of spread, severity, and inaction. *BMJ*, *368*, m1036. <https://doi.org/10.1136/bmj.m1036>
- Marmiroli, N., Peano, C., & Maestri, E. (2003). Advanced PCR techniques in identifying food components. *Food Authenticity and Traceability*, 3–33. <https://doi.org/10.1533/9781855737181.1.3>
- Muramatsu, T., Takemoto, C., Kim, Y.-T., Wang, H., Nishii, W., Terada, T., Shirouzu, M., & Yokoyama, S. (2016). SARS-CoV 3CL protease cleaves its C-terminal autoprocessing site by novel subsite cooperativity. *Proceedings of the National Academy of Sciences*, *113*(46), 12997–13002. <https://doi.org/10.1073/pnas.1601327113>
- Nanopores. (n.d.). *Types of nanopores*. Oxford Nanopore Technologies. <https://nanoporetech.com/how-it-works/types-of-nanopores>
- Nanoporetech. (n.d.-a). *Flow cells and nanopores*. Oxford Nanopore Technologies. <https://nanoporetech.com/how-it-works/flow-cells-and-nanopores>
- Nanoporetech. (n.d.-b). *How basecalling works*. Oxford Nanopore Technologies. <https://nanoporetech.com/how-it-works/basecalling>
- Nanoporetech. (2019). *How nanopore sequencing works*. Oxford Nanopore Technologies. <https://nanoporetech.com/how-it-works>
- Nanoporetech. (2021b). Ligation Sequencing Kit SQK-LSK109. Nanoporetech.com. <https://store.nanoporetech.com/ligation-sequencing-kit.html>
- Nanoporetech. (2021c). Rapid Barcoding Kit 96 SQK-RBK110.96. Nanoporetech.com. <https://store.nanoporetech.com/rapid-barcoding-kit-1.htm>

- Naqvi, A. A. T., Fatima, K., Mohammad, T., Fatima, U., Singh, I. K., Singh, A., Atif, S. M., Hariprasad, G., Hasan, G. M., & Hassan, Md. I. (2020). Insights into SARS-CoV-2 genome, structure, evolution, pathogenesis and therapies: Structural genomics approach. *Biochimica et Biophysica Acta. Molecular Basis of Disease*, 1866(10), 165878. <https://doi.org/10.1016/j.bbadis.2020.165878>
- Payne, S. (2017). Family Coronaviridae. *Viruses*, 149–158. <https://doi.org/10.1016/b978-0-12-803109-4.00017-9>
- Petrackova, A., Vasinek, M., Sedlarikova, L., Dyskova, T., Schneiderova, P., Novosad, T., Papajik, T., & Kriegova, E. (2019). Standardization of Sequencing Coverage Depth in NGS: Recommendation for Detection of Clonal and Subclonal Mutations in Cancer Diagnostics. *Frontiers in Oncology*, 9. <https://doi.org/10.3389/fonc.2019.00851>
- Quick, J. (2020). nCoV-2019 sequencing protocol v3 (LoCost). *Www.protocols.io*. <https://www.protocols.io/view/ncov-2019-sequencing-protocol-v3-locost-bh42j8ye>
- Rosenthal, S. H., Gerasimova, A., Ruiz-Vega, R., Livingston, K., Kagan, R. M., Liu, Y., Anderson, B., Owen, R., Bernstein, L., Smolgovsky, A., Xu, D., Chen, R., Grupe, A., Tanpaiboon, P., & Lacbawan, F. (2022). Development and validation of a high throughput SARS-CoV-2 whole genome sequencing workflow in a clinical laboratory. *Scientific Reports*, 12(1). <https://doi.org/10.1038/s41598-022-06091-0>
- Shen, C.-H. (2019, January 1). *Chapter 9 - Amplification of Nucleic Acids* (C.-H. Shen, Ed.). ScienceDirect; Academic Press. <https://www.sciencedirect.com/science/article/pii/B9780128028230000092>
- Shum, J., & Paul, N. (2009). Chemically modified primers for improved multiplex polymerase chain reaction. *Analytical Biochemistry*, 388(2), 266–272. <https://doi.org/10.1016/j.ab.2009.02.033>
- Sint, D., Raso, L., & Traugott, M. (2012). Advances in multiplex PCR: balancing primer efficiencies and improving detection success. *Methods in Ecology and Evolution*, 3(5), 898–905. <https://doi.org/10.1111/j.2041-210x.2012.00215.x>
- Thermo Fisher. (n.d.). *PCR Troubleshooting Guide - US*. *Www.thermofisher.com*. Retrieved May 30, 2022, from <https://www.thermofisher.com/id/en/home/life-science/cloning/cloning-learning-center/invitrogen-school-of-molecular-biology/pcr-education/pcr-reagents-enzymes/pcr-troubleshooting.html>
- Thermo Scientific. (n.d.). *General Recommendations for DNA Electrophoresis*. https://assets.thermofisher.com/TFS-Assets/LSG/manuals/MAN0012614_Gen_Recommend_DNA_Electrophoresis_UG.pdf
- Tyson, J. R., James, P., Stoddart, D., Sparks, N., Wickenhagen, A., Hall, G., Choi, J. H., Lapointe, H., Kamelian, K., Smith, A. D., Prystajacky, N., Goodfellow, I., Wilson, S. J., Harrigan, R., Snutch, T. P., Loman, N. J., & Quick, J. (2020). *Improvements to the ARTIC multiplex PCR method for SARS-CoV-2 genome sequencing using nanopore*. <https://doi.org/10.1101/2020.09.04.283077>
- Vernet, G. (2017). Genomics of Infectious Diseases and Private Industry. *Genetics and Evolution of Infectious Diseases*, 421–434. <https://doi.org/10.1016/b978-0-12-799942-5.00018-4>
- Vural-Ozdeniz, M., Akturk, A., Demirdizen, M., Leka, R., Acar, R., & Konu, O. (2021). CoVrimer: A tool for aligning SARS-CoV-2 primer sequences and selection of conserved/degenerate primers. *Genomics*, 113(5), 3174–3184. <https://doi.org/10.1016/j.ygeno.2021.07.020>
- WHO. (2020). *Origin of SARS-CoV-2*. https://apps.who.int/iris/bitstream/handle/10665/332197/WHO-2019-nCoV-FAQ-Virus_origin-2020.1-eng.pdf
- WHO. (2021). Genomic sequencing of SARS-CoV-2: a guide to implementation for maximum impact on public health, 8 January 2021. In *apps.who.int*. World Health Organization. <https://apps.who.int/iris/handle/10665/338480>
- Xiao, T., & Zhou, W. (2020). The third generation sequencing: the advanced approach to genetic diseases. *Translational Pediatrics*, 9(2), 163–173. <https://doi.org/10.21037/tp.2020.03.06>
- Yin, R., Kwok, C. K., & Zheng, J. (2019). Whole Genome Sequencing Analysis. *Encyclopedia of Bioinformatics and Computational Biology*, 176–183. <https://doi.org/10.1016/b978-0-12->

809633-8.20095-2

- Zhang, J., Lan, Y., & Sanyal, S. (2020). Membrane heist: Coronavirus host membrane remodeling during replication. *Biochimie*, *179*, 229–236. <https://doi.org/10.1016/j.biochi.2020.10.010>
- Zimmermann, F., Urban, M., Krüger, C., Walter, M., Wölfel, R., & Zwirgmaier, K. (2022). In vitro evaluation of the effect of mutations in primer binding sites on detection of SARS-CoV-2 by RT-qPCR. *Journal of Virological Methods*, *299*, 114352. <https://doi.org/10.1016/j.jviromet.2021.114352>

APPENDICES

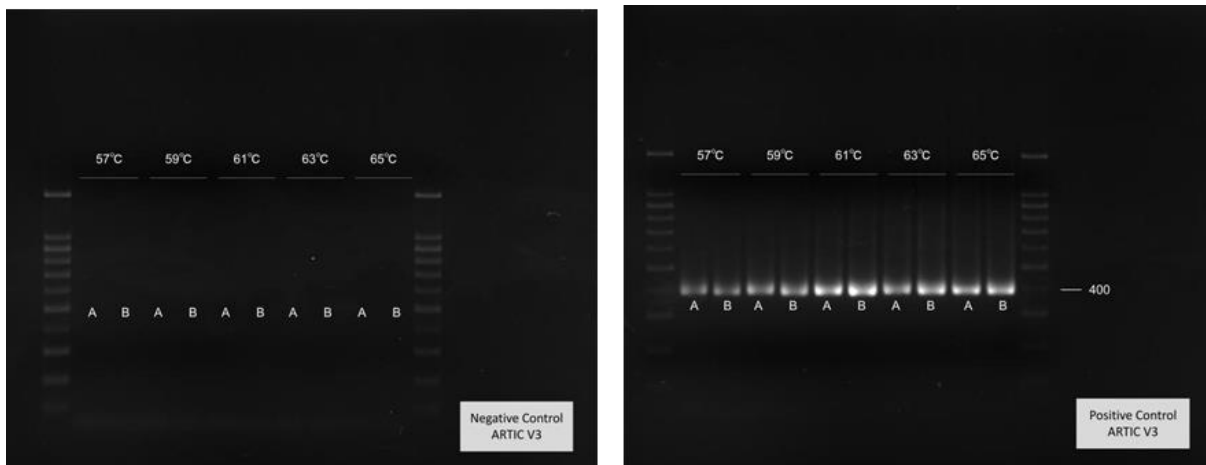
Supplementary Table 1. Nextflow WF-ARTIC Command Script

Nextflow WF-ARTIC Command Script

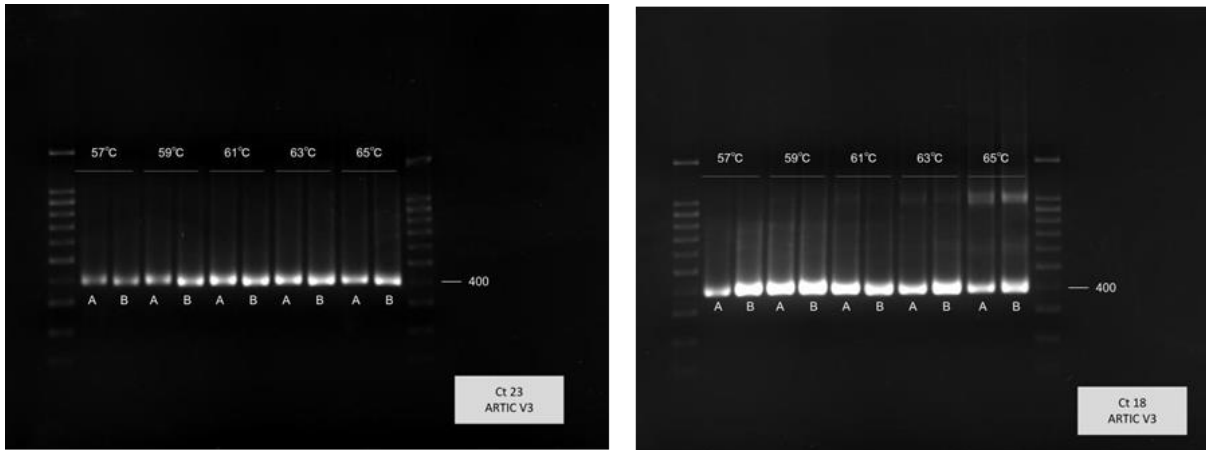
```
--fastq (*) path (to demultiplexed folder of FASTQ files)
--scheme_name SARS-CoV-2 --scheme_version V1200
--medaka_model r941_min_hac_variant_g507
    --the medaka model should reflect the basecaller used
    -r941_prom_variant_g360 is appropriate for all Guppy 4 runs
    -r941_min_hac_variant_g507 for HAC called Guppy 5.x data
    -r941_min_fast_variant_g507 for FAST called Guppy 5.x data
--out_dir my_result_folder
--profile standard (conda if docker unavailable)
```

Full script

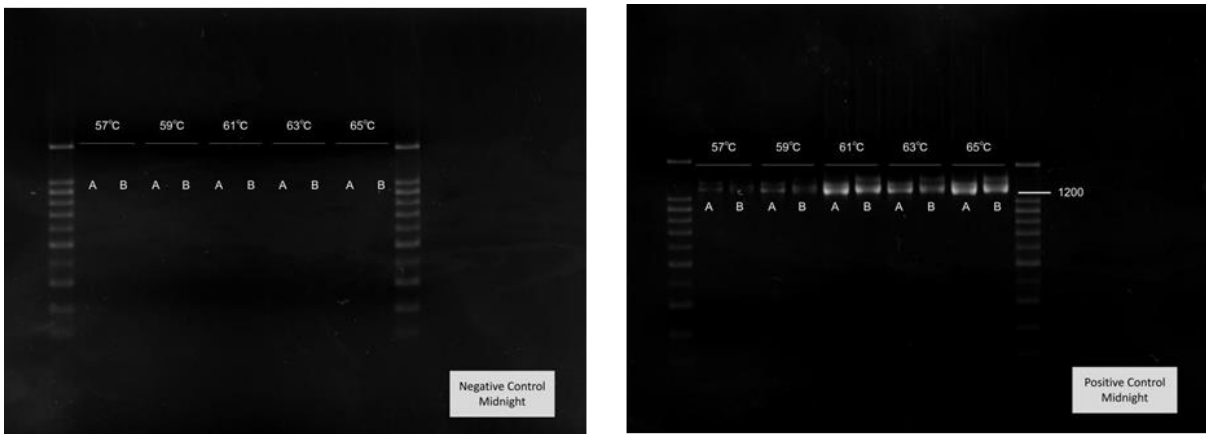
```
~/nextflow run ~/wf-artic-master --fastq FASTQ_PASS --output output --
medaka_model r941_min_hac_variant_g507 --scheme_version V1200
```



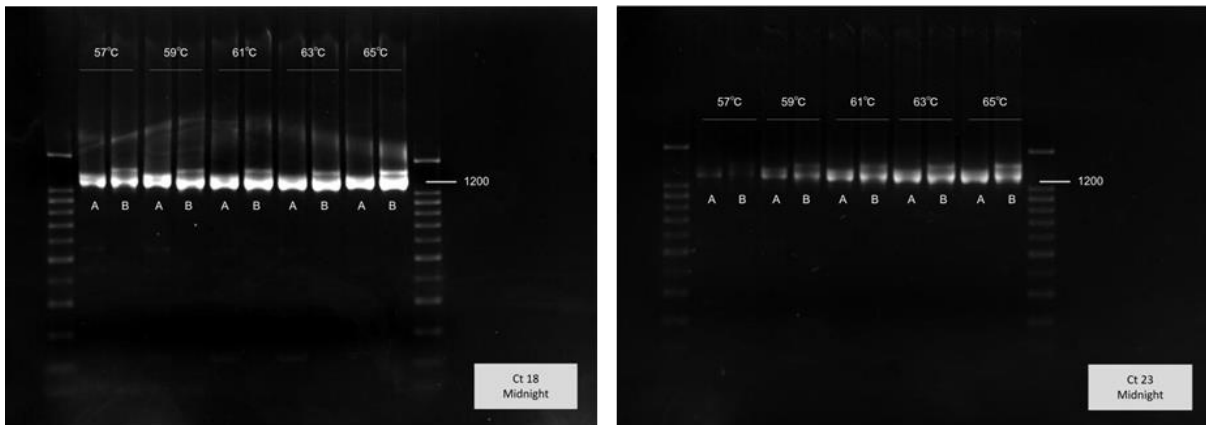
Supplementary Figure 1. Gel electrophoresis results from the negative control and positive control using ARTIC V3 primer.



Supplementary Figure 2. Gel electrophoresis result of Ct 18 and Ct 23 sample using ARTIC V3 primer.

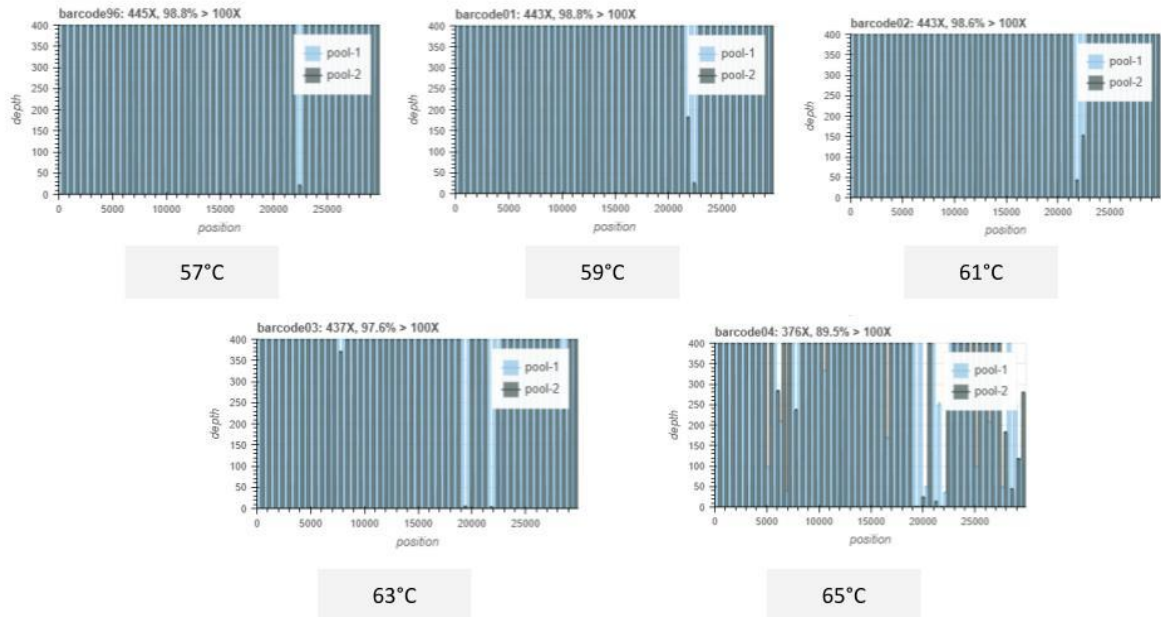


Supplementary Figure 3. Gel electrophoresis results from negative control and positive control using Midnight primer.



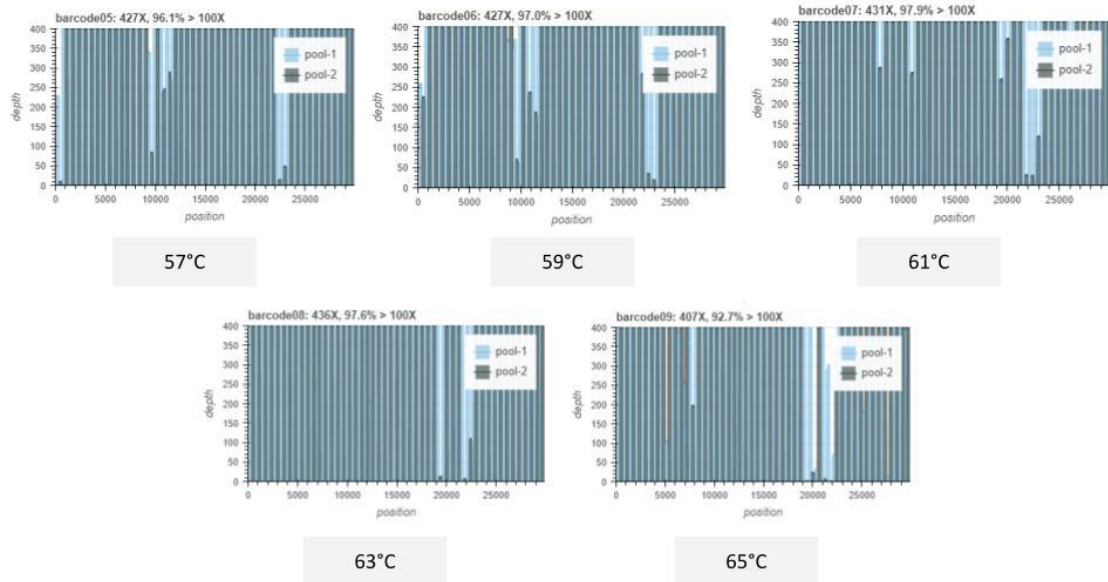
Supplementary Figure 4. Gel electrophoresis result of Ct 18 and Ct 23 using Midnight primer.

SQK-LSK109 // ARTIC V3 // Ct 18



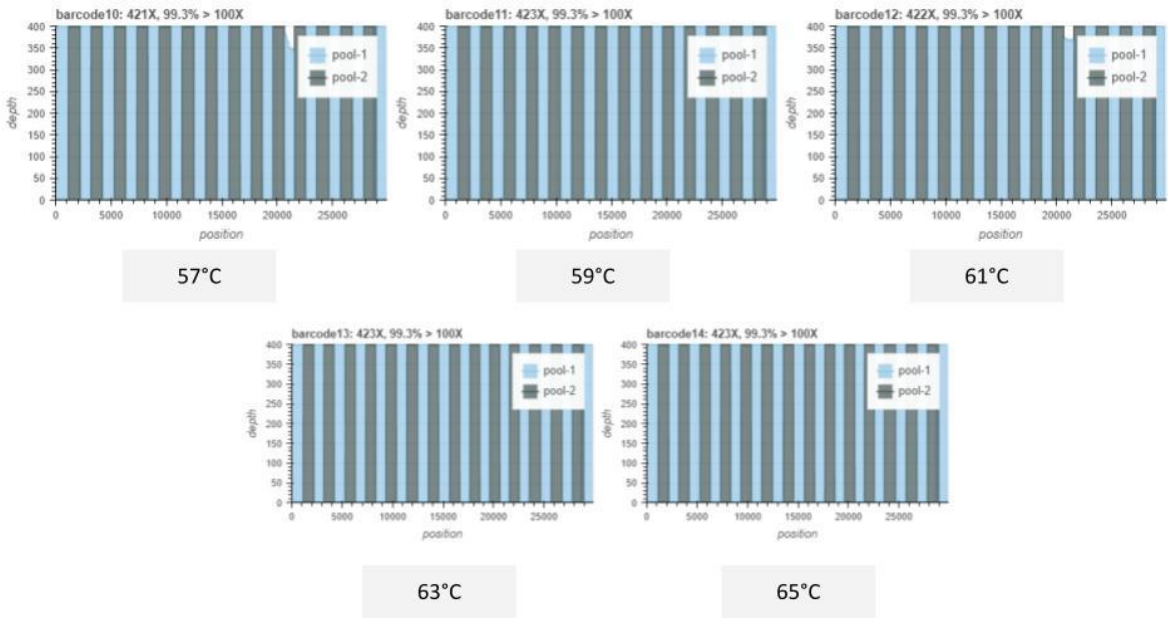
Supplementary Figure 5. Effect of different annealing temperatures on Ct 18 ARTIC V3 sequencing using SQK-LSK109 as seen on the amplicon pool.

SQK-LSK109 // ARTIC V3 // Ct 23



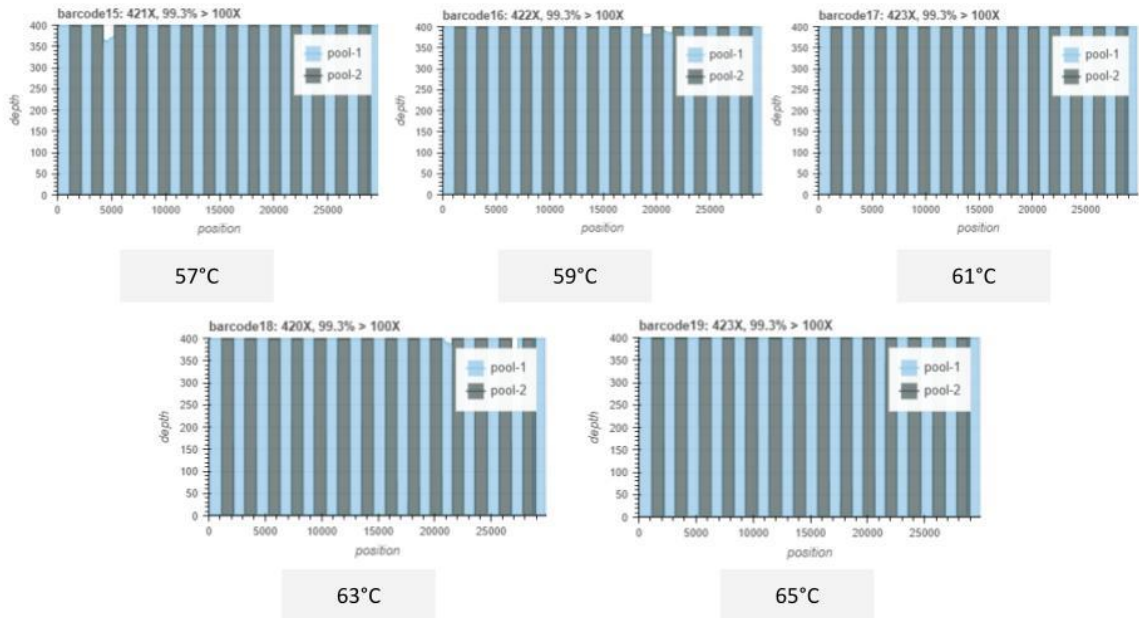
Supplementary Figure 6. Effect of different annealing temperatures on Ct 23 ARTIC V3 sequencing using SQK-LSK109 as seen on the amplicon pool.

SQK-LSK109 // MIDNIGHT // Ct 18



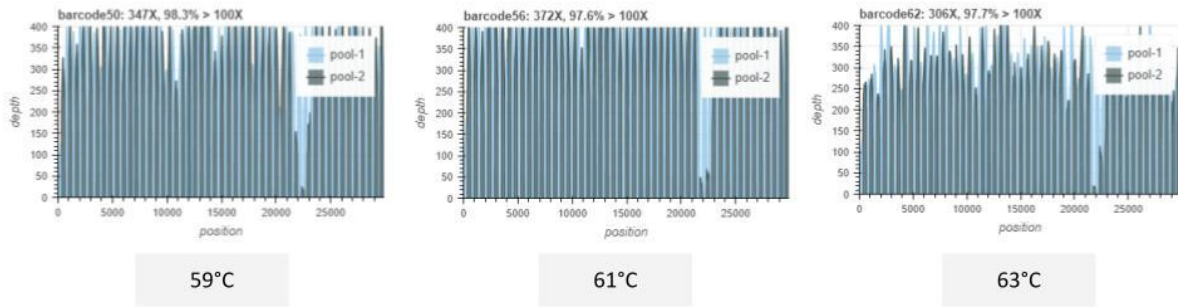
Supplementary Figure 7. Effect of different annealing temperatures on Ct 18 Midnight sequencing using SQK-LSK109 as seen on the amplicon pool.

SQK-LSK109 // MIDNIGHT // Ct 23



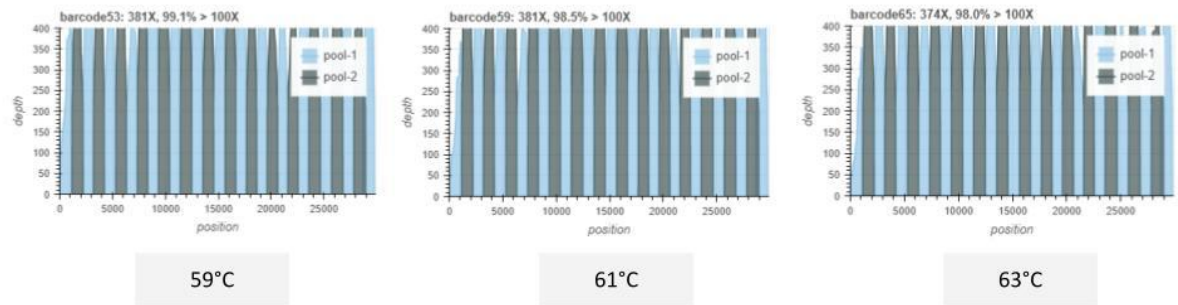
Supplementary Figure 8. Effect of different annealing temperatures on Ct 23 Midnight sequencing using SQK-LSK109 as seen on the amplicon pool.

SQK-RBK110.96 // ARTIC V3 // Ct 25

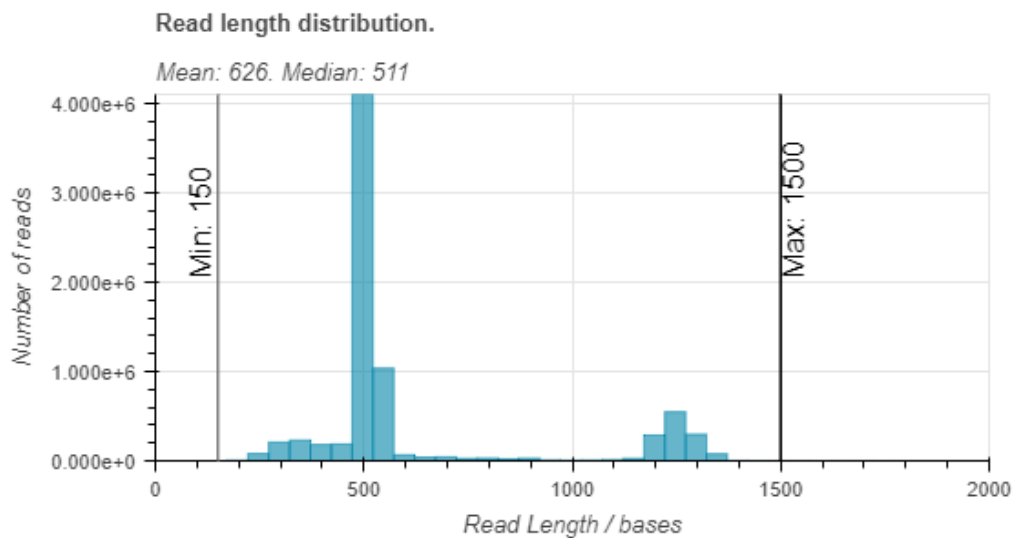


Supplementary Figure 9. Effect of different annealing temperatures on Ct 25 ARTIC V3 sequencing using SQK-RBK110.96 as seen on the amplicon pool.

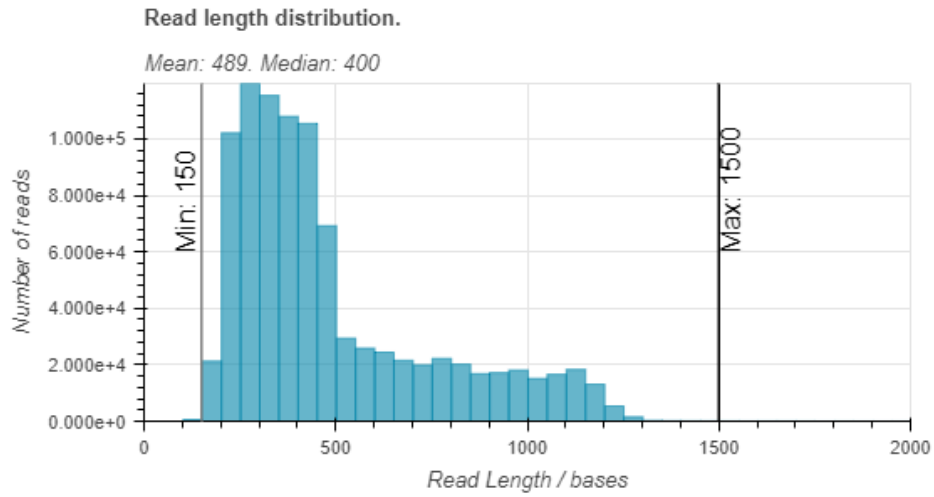
SQK-RBK110.96 // MIDNIGHT // Ct 25



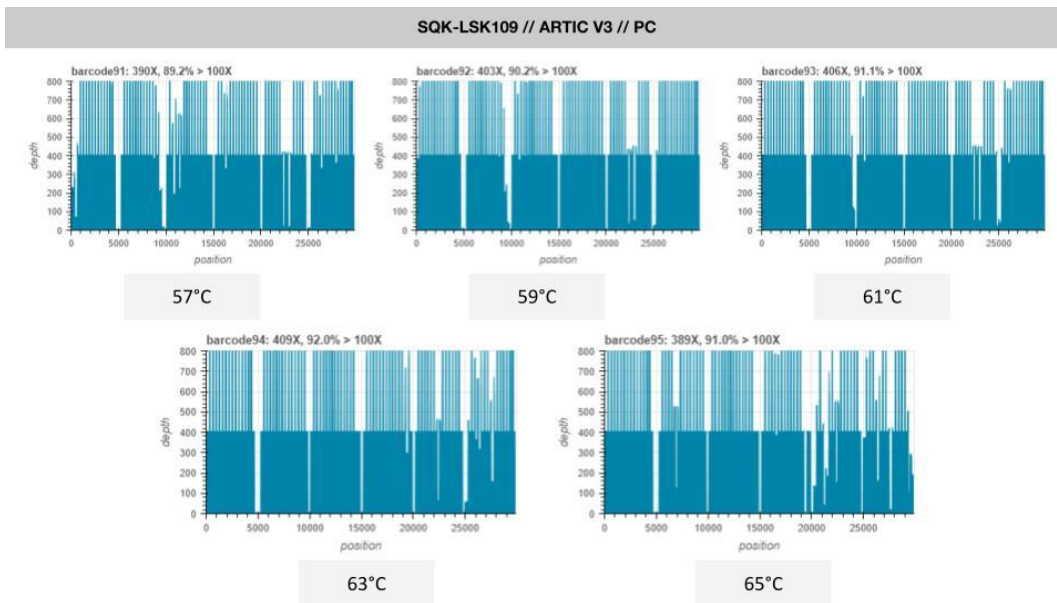
Supplementary Figure 10. Effect of different annealing temperatures on Ct 25 Midnight sequencing using SQK-RBK110.96 as seen on the amplicon pool.



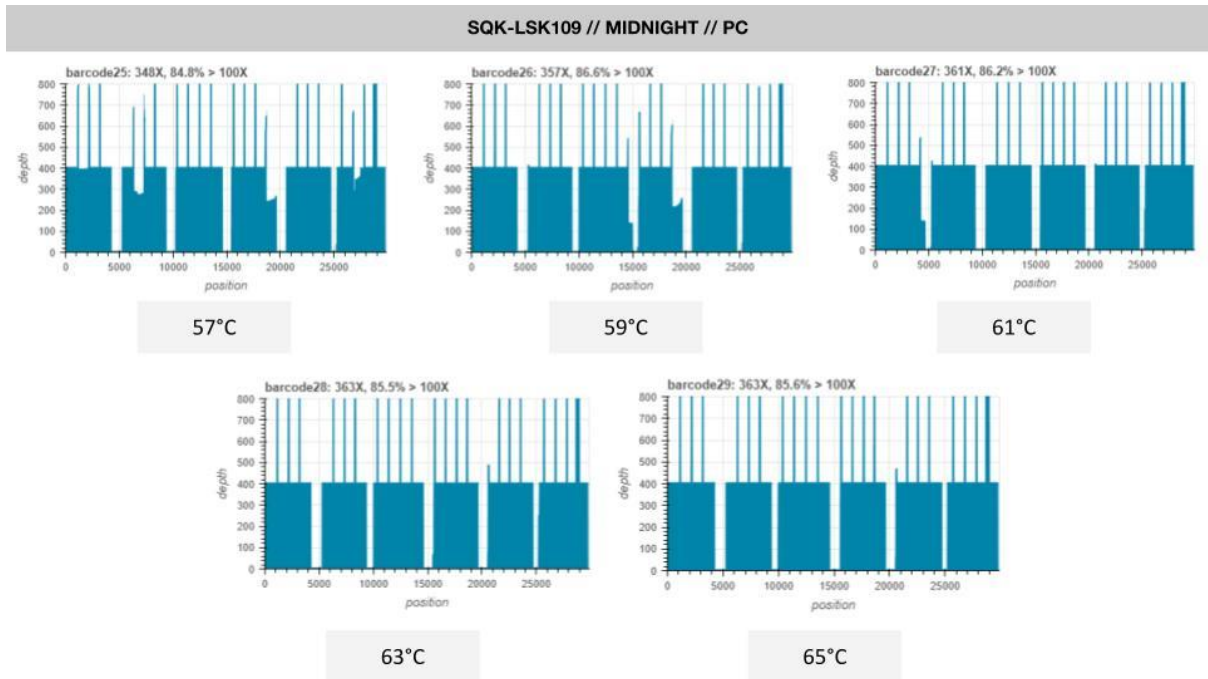
Supplementary Figure 11. Read length distribution of SQK-LSK109 sequencing.



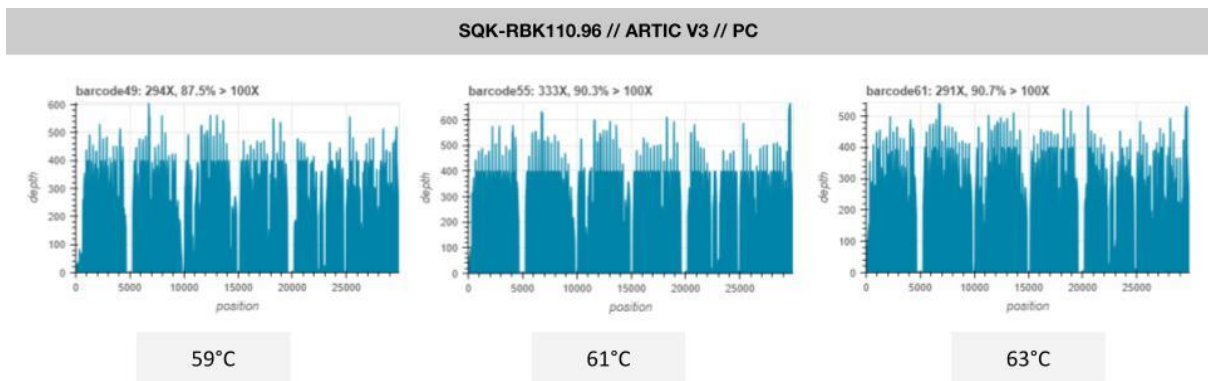
Supplementary Figure 12. Read length distribution of SQK-RBK110.96 sequencing.



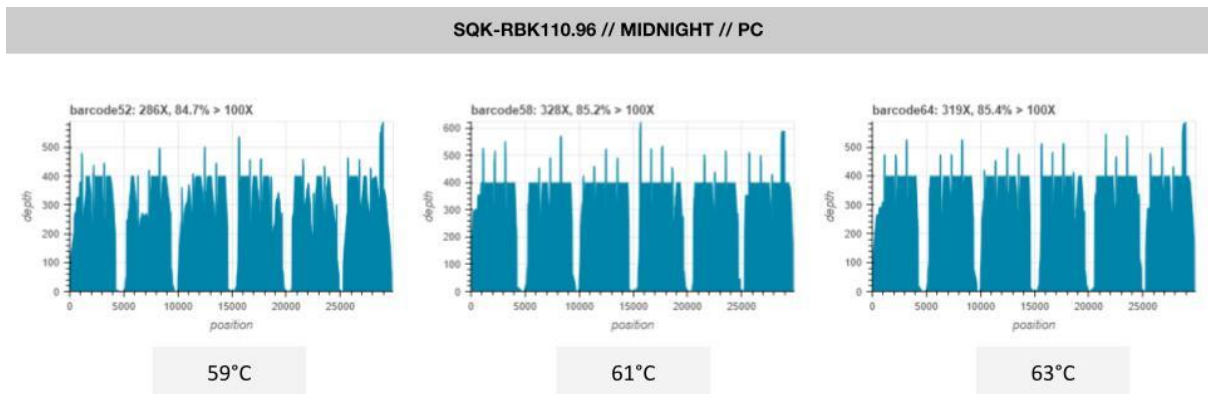
Supplementary Figure 13. Effect of different annealing temperatures on the positive control (PC) ARTIC V3 sequencing using SQK-LSK109.



Supplementary Figure 14. Effect of different annealing temperatures on the positive control (PC) Midnight sequencing using SQK-LSK109.



Supplementary Figure 15. Effect of different annealing temperatures on the positive control (PC) ARTIC V3 sequencing using SQK-RBK110.96.



Supplementary Figure 16. Effect of different annealing temperatures on the positive control (PC) Midnight sequencing using SQK-RBK110.96.

Supplementary Table 2. ARTIC V3 primer set (GitHub artic-ncov2019 V3, 2021).

PRIMER NAME	POOL	SEQUENCES	LENGTH (bp)	%GC	Tm
nCoV-2019_1_LEFT	nCoV-2019_1	ACCAACCAACTTTTCGATCTCTTGT	24	41.67	60.69
nCoV-2019_1_RIGHT	nCoV-2019_1	CATCTTTAAGATGTTGACGTGCCT C	25	44	60.45
nCoV-2019_2_LEFT	nCoV-2019_2	CTGTTTTACAGGTTCCGCGACGT	22	50	61.67
nCoV-2019_2_RIGHT	nCoV-2019_2	TAAGGATCAGTGCCAAGCTCGT	22	50	61.74
nCoV-2019_3_LEFT	nCoV-2019_1	CGGTAATAAAGGAGCTGGTGGC	22	54.55	61.32
nCoV-2019_3_RIGHT	nCoV-2019_1	AAGGTGTCTGCAATTCATAGCTCT	24	41.67	60.32
nCoV-2019_4_LEFT	nCoV-2019_2	GGTGTATACTGCTGCCGTGAAC	22	54.55	61.56
nCoV-2019_4_RIGHT	nCoV-2019_2	CACAAGTAGTGGCACCTTCTTTAG T	25	44	60.97
nCoV-2019_5_LEFT	nCoV-2019_1	TGGTGAAACTTCATGGCAGACG	22	50	61.39
nCoV-2019_5_RIGHT	nCoV-2019_1	ATTGATGTTGACTTTCTCTTTTTGG AGT	28	32.14	60.17
nCoV-2019_6_LEFT	nCoV-2019_2	GGTGTGTTGGAGAAGGTTCCG	22	54.55	61.64
nCoV-2019_6_RIGHT	nCoV-2019_2	TAGCGGCCTTCTGTAAACACG	22	50	61.18
nCoV-2019_7_LEFT	nCoV-2019_1	ATCAGAGGCTGCTCGTGTGTA	22	50	61.73
nCoV-2019_7_LEFT_alt0	nCoV-2019_1	CATTTGCATCAGAGGCTGCTCG	22	54.55	62.44
nCoV-2019_7_RIGHT	nCoV-2019_1	TGCACAGGTGACAATTTGTCCA	22	45.45	60.95
nCoV-2019_7_RIGHT_alt5	nCoV-2019_1	AGGTGACAATTTGTCCACCGAC	22	50	61.07
nCoV-2019_8_LEFT	nCoV-2019_2	AGAGTTTCTTAGAGACGGTTGGGA	24	45.83	61
nCoV-2019_8_RIGHT	nCoV-2019_2	GCTTCAACAGCTTCACTAGTAGGT	24	45.83	60.56
nCoV-2019_9_LEFT	nCoV-2019_1	TCCCACAGAAGTGTTAACAGAGGA	24	45.83	61.18
nCoV-2019_9_LEFT_alt4	nCoV-2019_1	TTCCCACAGAAGTGTTAACAGAGG	24	45.83	60.44

nCoV-2019_9_RIGHT	nCoV-2019_1	ATGACAGCATCTGCCACAACAC	22	50	61.71
nCoV-2019_9_RIGHT_alt2	nCoV-2019_1	GACAGCATCTGCCACAACACAG	22	54.55	62.26
nCoV-2019_10_LEFT	nCoV-2019_2	TGAGAAGTGCTCTGCCTATACAGT	24	45.83	61.12
nCoV-2019_10_RIGHT	nCoV-2019_2	TCATCTAACCAATCTTCTTCTTGCT CT	27	37.04	60.31
nCoV-2019_11_LEFT	nCoV-2019_1	GGAATTTGGTGCCACTTCTGCT	22	50	61.66
nCoV-2019_11_RIGHT	nCoV-2019_1	TCATCAGATTCAACTTGCATGGCA	24	41.67	61.35
nCoV-2019_12_LEFT	nCoV-2019_2	AAACATGGAGGAGGTGTTGCAG	22	50	61.08
nCoV-2019_12_RIGHT	nCoV-2019_2	TTCACTCTTCATTTCCAAAAAGCTT GA	27	33.33	60.36
nCoV-2019_13_LEFT	nCoV-2019_1	TCGCACAAATGTCTACTTAGCTGT	24	41.67	60.56
nCoV-2019_13_RIGHT	nCoV-2019_1	ACCACAGCAGTTAAACACCCT	22	45.45	60.36
nCoV-2019_14_LEFT	nCoV-2019_2	CATCCAGATTCTGCCACTCTTGT	23	47.83	60.62
nCoV-2019_14_LEFT_alt4	nCoV-2019_2	TGGCAATCTTCATCCAGATTCTGC	24	45.83	61.47
nCoV-2019_14_RIGHT	nCoV-2019_2	AGTTTCCACACAGACAGGCATT	22	45.45	60.42
nCoV-2019_14_RIGHT_alt2	nCoV-2019_2	TGCGTGTTTCTTCTGCATGTGC	22	50	62.76
nCoV-2019_15_LEFT	nCoV-2019_1	ACAGTGCTTAAAAAGTGTAAGT GCC	27	37.04	61.32
nCoV-2019_15_LEFT_alt1	nCoV-2019_1	AGTGCTTAAAAAGTGTAAGTGC CT	26	34.62	60.13
nCoV-2019_15_RIGHT	nCoV-2019_1	AACAGAACTGTAGCTGGCACT	22	45.45	60.16
nCoV-2019_15_RIGHT_alt3	nCoV-2019_1	ACTGTAGCTGGCACTTTGAGAGA	23	47.83	61.57
nCoV-2019_16_LEFT	nCoV-2019_2	AATTTGGAAGAAGCTGCTCGGT	22	45.45	60.82
nCoV-2019_16_RIGHT	nCoV-2019_2	CACAACCTGCGTGTGGAGGTTA	22	50	61.32

nCoV-2019_17_LEFT	nCoV-2019_1	CTTCTTTCTTTGAGAGAAGTGAGG ACT	27	40.74	60.69
nCoV-2019_17_RIGHT	nCoV-2019_1	TTTGTGGAGTGTTAACAATGCAG T	25	36	60.11
nCoV-2019_18_LEFT	nCoV-2019_2	TGGAAATACCCACAAGTTAATGGT TTAAC	29	34.48	60.69
nCoV-2019_18_LEFT_alt2	nCoV-2019_2	ACTTCTATTAATGGGCAGATAACA ACTGT	30	33.33	61.38
nCoV-2019_18_RIGHT	nCoV-2019_2	AGCTTGTTTACCACACGTACAAGG	24	45.83	61.51
nCoV-2019_18_RIGHT_alt1	nCoV-2019_2	GCTTGTTTACCACACGTACAAGG	23	47.83	60.3
nCoV-2019_19_LEFT	nCoV-2019_1	GCTGTTATGTACATGGGCACACT	23	47.83	61.18
nCoV-2019_19_RIGHT	nCoV-2019_1	TGTCCAACCTTAGGGTCAATTTCTGT	25	40	60.4
nCoV-2019_20_LEFT	nCoV-2019_2	ACAAAGAAAACAGTTACACAACAA CCA	27	33.33	60.68
nCoV-2019_20_RIGHT	nCoV-2019_2	ACGTGGCTTTATTAGTTGCATTGTT	25	36	60.28
nCoV-2019_21_LEFT	nCoV-2019_1	TGGCTATTGATTATAAACTACTACAC ACCC	29	37.93	61.49
nCoV-2019_21_LEFT_alt2	nCoV-2019_1	GGCTATTGATTATAAACTACTACACA CCCT	29	37.93	61.29
nCoV-2019_21_RIGHT	nCoV-2019_1	TAGATCTGTGTGGCCAACCTCT	22	50	60.83
nCoV-2019_21_RIGHT_alt0	nCoV-2019_1	GATCTGTGTGGCCAACCTCTTC	22	54.55	61.2
nCoV-2019_22_LEFT	nCoV-2019_2	ACTACCGAAGTTGTAGGAGACATT ATACT	29	37.93	61.25
nCoV-2019_22_RIGHT	nCoV-2019_2	ACAGTATTCTTTGCTATAGTAGTCG GC	27	40.74	60.73
nCoV-2019_23_LEFT	nCoV-2019_1	ACAACACTAACATAGTTACACGG TGT	27	37.04	60.26

nCoV-2019_23_RIGHT	nCoV-2019_1	ACCAGTACAGTAGGTTGCAATAGT G	25	44	60.57
nCoV-2019_24_LEFT	nCoV-2019_2	AGGCATGCCTTCTTACTGTACTG	23	47.83	60.37
nCoV-2019_24_RIGHT	nCoV-2019_2	ACATTCTAACCATAGCTGAAATCG GG	26	42.31	61.19
nCoV-2019_25_LEFT	nCoV-2019_1	GCAATTGTTTTTCAGCTATTTTGCA GT	27	33.33	60.73
nCoV-2019_25_RIGHT	nCoV-2019_1	ACTGTAGTGACAAGTCTCTCGCA	23	47.83	61.3
nCoV-2019_26_LEFT	nCoV-2019_2	TTGTGATACATTCTGTGCTGGTAG T	25	40	60.28
nCoV-2019_26_RIGHT	nCoV-2019_2	TCCGCACTATCACCAACATCAG	22	50	60.42
nCoV-2019_27_LEFT	nCoV-2019_1	ACTACAGTCAGCTTATGTGTCAAC C	25	44	60.8
nCoV-2019_27_RIGHT	nCoV-2019_1	AATACAAGCACCAAGGTCACGG	22	50	61.13
nCoV-2019_28_LEFT	nCoV-2019_2	ACATAGAAGTTACTGGCGATAGTT GT	26	38.46	60.13
nCoV-2019_28_RIGHT	nCoV-2019_2	TGTTTAGACATGACATGAACAGGT GT	26	38.46	60.91
nCoV-2019_29_LEFT	nCoV-2019_1	ACTTGTGTTCCTTTTTGTGCTGC	24	41.67	61.39
nCoV-2019_29_RIGHT	nCoV-2019_1	AGTGTACTCTATAAGTTTTGATGGT GTGT	29	34.48	60.69
nCoV-2019_30_LEFT	nCoV-2019_2	GCACAATAATGGTGACTTTTTGC A	25	40	61.19
nCoV-2019_30_RIGHT	nCoV-2019_2	ACCACTAGTAGATACACAAACACC AG	26	42.31	60.3
nCoV-2019_31_LEFT	nCoV-2019_1	TTCTGAGTACTGTAGGCACGGC	22	54.55	62.03
nCoV-2019_31_RIGHT	nCoV-2019_1	ACAGAATAAACACCAGGTAAGAAT GAGT	28	35.71	60.69

nCoV-2019_32_LEFT	nCoV-2019_2	TGGTGAATACAGTCATGTAGTTGC C	25	44	61.09
nCoV-2019_32_RIGHT	nCoV-2019_2	AGCACATCACTACGCAACTTTAGA	24	41.67	60.56
nCoV-2019_33_LEFT	nCoV-2019_1	ACTTTTGAAGAAGCTGCGCTGT	22	45.45	61.58
nCoV-2019_33_RIGHT	nCoV-2019_1	TGGACAGTAAACTACGTCATCAAG C	25	44	61.08
nCoV-2019_34_LEFT	nCoV-2019_2	TCCCATCTGGTAAAGTTGAGGGT	23	47.83	61.02
nCoV-2019_34_RIGHT	nCoV-2019_2	AGTGAAATTGGGCCTCATAGCA	22	45.45	60.03
nCoV-2019_35_LEFT	nCoV-2019_1	TGTTTCGCATTCAACCAGGACAG	22	50	61.39
nCoV-2019_35_RIGHT	nCoV-2019_1	ACTTCATAGCCACAAGGTTAAAGT CA	26	38.46	60.69
nCoV-2019_36_LEFT	nCoV-2019_2	TTAGCTTGGTTGTACGCTGCTG	22	50	61.44
nCoV-2019_36_RIGHT	nCoV-2019_2	GAACAAAGACCATTGAGTACTCTG GA	26	42.31	60.74
nCoV-2019_37_LEFT	nCoV-2019_1	ACACACCACTGGTTGTTACTCAC	23	47.83	60.93
nCoV-2019_37_RIGHT	nCoV-2019_1	GTCCACACTCTCCTAGCACCAT	22	54.55	61.48
nCoV-2019_38_LEFT	nCoV-2019_2	ACTGTGTTATGTATGCATCAGCTG T	25	40	60.86
nCoV-2019_38_RIGHT	nCoV-2019_2	CACCAAGAGTCAGTCTAAAGTAGC G	25	48	61.13
nCoV-2019_39_LEFT	nCoV-2019_1	AGTATTGCCCTATTTTCTTCATAAC TGTT	29	34.48	61
nCoV-2019_39_RIGHT	nCoV-2019_1	TGTAAGTGGACACATTGAGCCC	22	50	60.55
nCoV-2019_40_LEFT	nCoV-2019_2	TGCACATCAGTAGTCTTACTCTCA GT	26	42.31	61.25
nCoV-2019_40_RIGHT	nCoV-2019_2	CATGGCTGCATCACGGTCAAAT	22	50	62.09
nCoV-2019_41_LEFT	nCoV-2019_1	GTTCCCTTCCATCATATGCAGCT	23	47.83	60.75
nCoV-2019_41_RIGHT	nCoV-2019_1	TGGTATGACAACCATTAGTTTGGC T	25	40	60.75

nCoV-2019_42_LEFT	nCoV-2019_2	TGCAAGAGATGGTTGTGTTCCC	22	50	61.08
nCoV-2019_42_RIGHT	nCoV-2019_2	CCTACCTCCCTTTGTTGTGTTGT	23	47.83	60.69
nCoV-2019_43_LEFT	nCoV-2019_1	TACGACAGATGTCTTGTGCTGC	22	50	60.93
nCoV-2019_43_RIGHT	nCoV-2019_1	AGCAGCATCTACAGCAAAAGCA	22	45.45	61.14
nCoV-2019_44_LEFT	nCoV-2019_2	TGCCACAGTACGTCTACAAGCT	22	50	61.66
nCoV-2019_44_LEFT_alt3	nCoV-2019_2	CCACAGTACGTCTACAAGCTGG	22	54.55	60.67
nCoV-2019_44_RIGHT	nCoV-2019_2	AACCTTTCCACATACCGCAGAC	22	50	60.87
nCoV-2019_44_RIGHT_alt0	nCoV-2019_2	CGCAGACGGTACAGACTGTGTT	22	54.55	62.77
nCoV-2019_45_LEFT	nCoV-2019_1	TACCTACAACCTTGTGCTAATGACC C	25	44	60.57
nCoV-2019_45_LEFT_alt2	nCoV-2019_1	AGTATGTACAAATACCTACAACCTTG TGCT	29	34.48	60.94
nCoV-2019_45_RIGHT	nCoV-2019_1	AAATTGTTTCTTCATGTTGGTAGTT AGAGA	30	30	60.01
nCoV-2019_45_RIGHT_alt7	nCoV-2019_1	TTCATGTTGGTAGTTAGAGAAAGT GTGTC	29	37.93	61.53
nCoV-2019_46_LEFT	nCoV-2019_2	TGTCGCTTCCAAGAAAAGGACG	22	50	61.38
nCoV-2019_46_LEFT_alt1	nCoV-2019_2	CGCTTCCAAGAAAAGGACGAAGA	23	47.83	61.35
nCoV-2019_46_RIGHT	nCoV-2019_2	CACGTTACCTAAGTTGGCGTA	22	50	60.86
nCoV-2019_46_RIGHT_alt2	nCoV-2019_2	CACGTTACCTAAGTTGGCGTAT	23	47.83	61.17
nCoV-2019_47_LEFT	nCoV-2019_1	AGGACTGGTATGATTTTGTAGAAA ACCC	28	39.29	61.42
nCoV-2019_47_RIGHT	nCoV-2019_1	AATAACGGTCAAAGAGTTTTAACCT CTC	28	35.71	60.06

nCoV-2019_48_LEFT	nCoV-2019_2	TGTTGACACTGACTTAACAAAGCC T	25	40	61.09
nCoV-2019_48_RIGHT	nCoV-2019_2	TAGATTACCAGAAGCAGCGTGC	22	50	60.74
nCoV-2019_49_LEFT	nCoV-2019_1	AGGAATTACTTGTGTATGCTGCTG A	25	40	60.57
nCoV-2019_49_RIGHT	nCoV-2019_1	TGACGATGACTTGGTTAGCATTAA TACA	28	35.71	61.05
nCoV-2019_50_LEFT	nCoV-2019_2	GTTGATAAGTACTTTGATTGTTACG ATGGT	30	33.33	60.59
nCoV-2019_50_RIGHT	nCoV-2019_2	TAACATGTTGTGCCAACCA	22	45.45	60.95
nCoV-2019_51_LEFT	nCoV-2019_1	TCAATAGCCGCCACTAGAGGAG	22	54.55	61.34
nCoV-2019_51_RIGHT	nCoV-2019_1	AGTGCATTAACATTGGCCGTGA	22	45.45	61.14
nCoV-2019_52_LEFT	nCoV-2019_2	CATCAGGAGATGCCACAACCTGC	22	54.55	61.83
nCoV-2019_52_RIGHT	nCoV-2019_2	GTTGAGAGCAAATTCATGAGGTC C	25	44	60.62
nCoV-2019_53_LEFT	nCoV-2019_1	AGCAAATGTTGGACTGAGACTGA	24	41.67	60.69
nCoV-2019_53_RIGHT	nCoV-2019_1	AGCCTCATAAACTCAGGTTCCC	23	47.83	60.31
nCoV-2019_54_LEFT	nCoV-2019_2	TGAGTTAACAGGACACATGTTAGA CA	26	38.46	60.18
nCoV-2019_54_RIGHT	nCoV-2019_2	AACCAAAAACCTTGTCCATTAGCAC A	25	36	60.11
nCoV-2019_55_LEFT	nCoV-2019_1	ACTCAACTTTACTTAGGAGGTATG AGCT	28	39.29	61.43
nCoV-2019_55_RIGHT	nCoV-2019_1	GGTGTACTCTCCTATTTGTACTTTA CTGT	29	37.93	60.54
nCoV-2019_56_LEFT	nCoV-2019_2	ACCTAGACCACCACTTAACCGA	22	50	60.49
nCoV-2019_56_RIGHT	nCoV-2019_2	ACACTATGCGAGCAGAAGGGTA	22	50	61.21
nCoV-2019_57_LEFT	nCoV-2019_1	ATTCTACACTCCAGGGACCACC	22	54.55	61.16
nCoV-2019_57_RIGHT	nCoV-2019_1	GTAATTGAGCAGGGTCGCCAAT	22	50	61.26

nCoV-2019_58_LEFT	nCoV-2019_2	TGATTTGAGTGTTGTCAATGCCAG A	25	40	61.44
nCoV-2019_58_RIGHT	nCoV-2019_2	CTTTTCTCCAAGCAGGGTTACGT	23	47.83	61.06
nCoV-2019_59_LEFT	nCoV-2019_1	TCACGCATGATGTTTCATCTGCA	23	43.48	61.42
nCoV-2019_59_RIGHT	nCoV-2019_1	AAGAGTCCTGTTACATTTTCAGCTT G	26	38.46	60.02
nCoV-2019_60_LEFT	nCoV-2019_2	TGATAGAGACCTTTATGACAAGTT GCA	27	37.04	60.53
nCoV-2019_60_RIGHT	nCoV-2019_2	GGTACCAACAGCTTCTCTAGTAGC	24	50	60.44
nCoV-2019_61_LEFT	nCoV-2019_1	TGTTTATCACCCGCGAAGAAGC	22	50	61.5
nCoV-2019_61_RIGHT	nCoV-2019_1	ATCACATAGACAACAGGTGCGC	22	50	61.25
nCoV-2019_62_LEFT	nCoV-2019_2	GGCACATGGCTTTGAGTTGACA	22	50	61.91
nCoV-2019_62_RIGHT	nCoV-2019_2	GTTGAACCTTTCTACAAGCCGC	22	50	60.35
nCoV-2019_63_LEFT	nCoV-2019_1	TGTTAAGCGTGTTGACTGGACT	22	45.45	60.16
nCoV-2019_63_RIGHT	nCoV-2019_1	ACAAACTGCCACCATCACAACC	22	50	61.85
nCoV-2019_64_LEFT	nCoV-2019_2	TCGATAGATATCCTGCTAATTCCAT TGT	28	35.71	60.11
nCoV-2019_64_RIGHT	nCoV-2019_2	AGTCTTGTAAGAGTGTCCAGAGG T	25	40	60.1
nCoV-2019_65_LEFT	nCoV-2019_1	GCTGGCTTTAGCTTGTGGGTTT	22	50	61.92
nCoV-2019_65_RIGHT	nCoV-2019_1	TGTCAGTCATAGAACAAACACCAA TAGT	28	35.71	60.9
nCoV-2019_66_LEFT	nCoV-2019_2	GGGTGTGGACATTGCTGCTAAT	22	50	61.21
nCoV-2019_66_RIGHT	nCoV-2019_2	TCAATTTCCATTTGACTCCTGGGT	24	41.67	60.45
nCoV-2019_67_LEFT	nCoV-2019_1	GTTGTCCAACAATTACCTGAAACTT ACT	28	35.71	60.43
nCoV-2019_67_RIGHT	nCoV-2019_1	CAACCTTAGAACTACAGATAAATC TTGGG	30	36.67	60.4

nCoV-2019_68_LEFT	nCoV-2019_2	ACAGGTTTCATCTAAGTGTGTGTGT	24	41.67	60.14
nCoV-2019_68_RIGHT	nCoV-2019_2	CTCCTTTATCAGAACCAGCACCA	23	47.83	60.31
nCoV-2019_69_LEFT	nCoV-2019_1	TGTCGCAAAATATACTCAACTGTGT CA	27	37.04	61.43
nCoV-2019_69_RIGHT	nCoV-2019_1	TCTTTATAGCCACGGAACCTCCA	23	47.83	61.14
nCoV-2019_70_LEFT	nCoV-2019_2	ACAAAAGAAAATGACTCTAAAGAG GGTTT	29	31.03	60.13
nCoV-2019_70_RIGHT	nCoV-2019_2	TGACCTTCTTTTAAAGACATAACAG CAG	28	35.71	60.27
nCoV-2019_71_LEFT	nCoV-2019_1	ACAAATCCAATTCAGTTGTCTTCCT ATTC	29	34.48	60.54
nCoV-2019_71_RIGHT	nCoV-2019_1	TGAAAAGAAAGGTAAGAACAAGT CCT	27	37.04	60.8
nCoV-2019_72_LEFT	nCoV-2019_2	ACACGTGGTGTTTATTACCCTGAC	24	45.83	61.04
nCoV-2019_72_RIGHT	nCoV-2019_2	ACTCTGAACTCACTTCCATCCAAC	25	44	60.97
nCoV-2019_73_LEFT	nCoV-2019_1	CAATTTTGTAATGATCCATTTTGG GTGT	29	31.03	60.29
nCoV-2019_73_RIGHT	nCoV-2019_1	CACCAGCTGTCCAACCTGAAGA	22	54.55	62.45
nCoV-2019_74_LEFT	nCoV-2019_2	ACATCACTAGGTTTCAAACCTTACT TGC	28	35.71	60.68
nCoV-2019_74_RIGHT	nCoV-2019_2	GCAACACAGTTGCTGATTCTCTTC	24	45.83	60.85
nCoV-2019_75_LEFT	nCoV-2019_1	AGAGTCCAACCAACAGAATCTATT GT	26	38.46	60.24
nCoV-2019_75_RIGHT	nCoV-2019_1	ACCACCAACCTTAGAATCAAGATT GT	26	38.46	60.69
nCoV-2019_76_LEFT	nCoV-2019_2	AGGGCAAACCTGGAAAGATTGCT	22	45.45	60.76
nCoV-2019_76_LEFT_alt3	nCoV-2019_2	GGGCAAACCTGGAAAGATTGCTGA	23	47.83	61.87
nCoV-2019_76_RIGHT	nCoV-2019_2	ACACCTGTGCCTGTAAACCAT	22	45.45	60.42

nCoV-2019_76_RIGHT_alt0	nCoV-2019_2	ACCTGTGCCTGTAAACCATTGA	23	43.48	60.69
nCoV-2019_77_LEFT	nCoV-2019_1	CCAGCAACTGTTTGTGGACCTA	22	50	60.75
nCoV-2019_77_RIGHT	nCoV-2019_1	CAGCCCCTATTAACAGCCTGC	22	54.55	61.59
nCoV-2019_78_LEFT	nCoV-2019_2	CAACTTACTCCTACTTGGCGTGT	23	47.83	60.55
nCoV-2019_78_RIGHT	nCoV-2019_2	TGTGTACAAAACTGCCATATTGC A	25	36	60.22
nCoV-2019_79_LEFT	nCoV-2019_1	GTGGTGATTCAACTGAATGCAGC	23	47.83	60.92
nCoV-2019_79_RIGHT	nCoV-2019_1	CATTCATCTGTGAGCAAAGGTGG	24	45.83	60.62
nCoV-2019_80_LEFT	nCoV-2019_2	TTGCCTTGGTGATATTGCTGCT	22	45.45	60.89
nCoV-2019_80_RIGHT	nCoV-2019_2	TGGAGCTAAGTTGTTAACAAGCG	24	41.67	60.02
nCoV-2019_81_LEFT	nCoV-2019_1	GCACTTGGAAAACCTCAAGATGTG G	25	44	61.24
nCoV-2019_81_RIGHT	nCoV-2019_1	GTGAAGTTCTTTTCTTGTGCAGGG	24	45.83	60.73
nCoV-2019_82_LEFT	nCoV-2019_2	GGGCTATCATCTTATGTCCTTCCC T	25	48	61.52
nCoV-2019_82_RIGHT	nCoV-2019_2	TGCCAGAGATGTCACCTAAATCAA	24	41.67	60.02
nCoV-2019_83_LEFT	nCoV-2019_1	TCCTTTGCAACCTGAATTAGACTCA	25	40	60.46
nCoV-2019_83_RIGHT	nCoV-2019_1	TTTGACTCCTTTGAGCACTGGC	22	50	61.33
nCoV-2019_84_LEFT	nCoV-2019_2	TGCTGTAGTTGTCTCAAGGGCT	22	50	61.61
nCoV-2019_84_RIGHT	nCoV-2019_2	AGGTGTGAGTAACTGTTACAAAC AAC	27	37.04	60.36
nCoV-2019_85_LEFT	nCoV-2019_1	ACTAGCACTCTCCAAGGGTGTT	22	50	61.03
nCoV-2019_85_RIGHT	nCoV-2019_1	ACACAGTCTTTTACTCCAGATTCCC	25	44	60.51
nCoV-2019_86_LEFT	nCoV-2019_2	TCAGGTGATGGCACAACAAGTC	22	50	61.07
nCoV-2019_86_RIGHT	nCoV-2019_2	ACGAAAGCAAGAAAAAGAAGTACG C	25	40	61.01
nCoV-2019_87_LEFT	nCoV-2019_1	CGACTACTAGCGTGCCTTTGTA	22	50	60.16

nCoV-2019_87_RIGHT	nCoV-2019_1	ACTAGGTTCCATTGTTCAAGGAGC	24	45.83	60.81
nCoV-2019_88_LEFT	nCoV-2019_2	CCATGGCAGATTCCAACGGTAC	22	54.55	61.58
nCoV-2019_88_RIGHT	nCoV-2019_2	TGGTCAGAATAGTGCCATGGAGT	23	47.83	61.4
nCoV-2019_89_LEFT	nCoV-2019_1	GTACGCGTTCCATGTGGTCATT	22	50	61.5
nCoV-2019_89_LEFT_alt2	nCoV-2019_1	CGCGTTCCATGTGGTCATTCAA	22	50	62.01
nCoV-2019_89_RIGHT	nCoV-2019_1	ACCTGAAAGTCAACGAGATGAAAC A	25	40	60.91
nCoV-2019_89_RIGHT_alt4	nCoV-2019_1	ACGAGATGAAACATCTGTTGTAC T	25	40	60.74
nCoV-2019_90_LEFT	nCoV-2019_2	ACACAGACCATTCCAGTAGCAGT	23	47.83	61.58
nCoV-2019_90_RIGHT	nCoV-2019_2	TGAAATGGTGAATTGCCCTCGT	22	45.45	60.82
nCoV-2019_91_LEFT	nCoV-2019_1	TCACTACCAAGAGTGTGTTAGAGG T	25	44	60.93
nCoV-2019_91_RIGHT	nCoV-2019_1	TTCAAGTGAGAACCAAAAGATAAT AAGCA	29	31.03	60.03
nCoV-2019_92_LEFT	nCoV-2019_2	TTTGTGCTTTTTAGCCTTTCTGCT	24	37.5	60.14
nCoV-2019_92_RIGHT	nCoV-2019_2	AGGTTCTGGCAATTAATTGTAAAA GG	27	37.04	60.53
nCoV-2019_93_LEFT	nCoV-2019_1	TGAGGCTGTTCTAAATCACCCA	23	47.83	61.59
nCoV-2019_93_RIGHT	nCoV-2019_1	AGGTCTTCCTTGCCATGTTGAG	22	50	60.55
nCoV-2019_94_LEFT	nCoV-2019_2	GGCCCCAAGGTTTACCCAATAA	22	50	60.56
nCoV-2019_94_RIGHT	nCoV-2019_2	TTTGGCAATGTTGTTCCCTTGAGG	23	43.48	60.18
nCoV-2019_95_LEFT	nCoV-2019_1	TGAGGGAGCCTTGAATACACCA	22	50	61.1
nCoV-2019_95_RIGHT	nCoV-2019_1	CAGTACGTTTTTGGCGAGGCTT	22	50	61.95
nCoV-2019_96_LEFT	nCoV-2019_2	GCCAACAACAACAAGGCCAAAC	22	50	61.82
nCoV-2019_96_RIGHT	nCoV-2019_2	TAGGCTCTGTTGGTGGGAATGT	22	50	61.36

nCoV-2019_97_LEFT	nCoV-2019_1	TGGATGACAAAGATCCAAATTTCA AAGA	28	32.14	60.22
nCoV-2019_97_RIGHT	nCoV-2019_1	ACACACTGATTAAGATTGCTATGT GAG	28	35.71	60.17
nCoV-2019_98_LEFT	nCoV-2019_2	AACAATTGCAACAATCCATGAGCA	24	37.5	60.5
nCoV-2019_98_RIGHT	nCoV-2019_2	TTCTCCTAAGAAGCTATTAATAATCA CATGG	30	33.33	60.01

Supplementary Table 3. Midnight primer set (Freed et al., 2020).

Primer Name	Sequence	Pool	Length	GC%	Tm
SARSCoV_1200_1_LEFT	ACCAACCAACTTTCGATCTCTTG T	1	24	41.67	60.69
SARSCoV_1200_1_RIGHT	GGTTGCATTCATTTGGTGACGC	1	22	50	61.49
SARSCoV_1200_3_LEFT	GGCTTGAAGAGAAGTTTAAGG AAGGT	1	26	42.31	61.19
SARSCoV_1200_3_RIGHT	GATTGTCCTCACTGCCGTCTTG	1	22	54.55	61.5
SARSCoV_1200_5_LEFT	ACCTACTAAAAGGCTGGTGG C	1	22	50	60.55
SARSCoV_1200_5_RIGHT	AGCATCTTGTAGAGCAGGTGG A	1	22	50	61.16
SARSCoV_1200_7_LEFT	ACCTGGTGTATACGTTGTCTTT GG	1	24	45.83	60.8
SARSCoV_1200_7_RIGHT	GCTGAAATCGGGGCCATTTGTA	1	22	50	61.53
SARSCoV_1200_9_LEFT	AGAAGTTACTGGCGATAGTTGT AATAACT	1	29	34.48	60.59
SARSCoV_1200_9_RIGHT	TGCTGATATGTCCAAAGCACCA	1	22	45.45	60.29
SARSCoV_1200_11_LEFT	AGACACCTAAGTATAAGTTTGT TCGCA	1	27	37.04	60.74
SARSCoV_1200_11_RIGHT	GCCACATGGAAATGGCTTGAT	1	22	50	61.8
SARSCoV_1200_13_LEFT	ACCTCTTACAACAGCAGCCAAA C	1	23	47.83	61.55

SARSCoV_1200_13_RIGHT	CGTCCTTTTCTTGGAAGCGACA	1	22	50	61.38
SARSCoV_1200_15_LEFT	TTTTAAGGAATTACTTGTGTAT GCTGCT	1	28	32.14	60.06
SARSCoV_1200_15_RIGHT	ACACACAACAGCATCGTCAGA G	1	22	50	61.12
SARSCoV_1200_17_LEFT	TCAAGCTTTTTGCAGCAGAAAC G	1	23	43.48	61.28
SARSCoV_1200_17_RIGHT	CCAAGCAGGGTTACGTGTAAG G	1	22	54.55	61.19
SARSCoV_1200_19_LEFT	GGCACATGGCTTTGAGTTGACA	1	22	50	61.91
SARSCoV_1200_19_RIGHT	CCTGTTGTCCATCAAAGTGTCC C	1	23	52.17	61.62
SARSCoV_1200_21_LEFT	TCTGTAGTTTCTAAGGTTGTCA AAGTGA	1	28	35.71	60.58
SARSCoV_1200_21_RIGHT	GCAGGGGGTAATTGAGTTCTG G	1	22	54.55	60.95
SARSCoV_1200_23_LEFT	ACTTTAGAGTCCAACCAACAGA ATCT	1	26	38.46	60.18
SARSCoV_1200_23_RIGHT	TGACTAGCTACACTACGTGCC	1	22	54.55	61.52
SARSCoV_1200_25_LEFT	TGCTGCTACTAAAATGTCAGAG TGT	1	25	40	60.51
SARSCoV_1200_25_RIGHT	CATTTCCAGCAAAGCCAAAGCC	1	22	50	61.45
SARSCoV_1200_27_LEFT	TGGATCACCGGTGGAATTGCTA	1	22	50	61.75
SARSCoV_1200_27_RIGHT	TGTTCGTTTAGGCGTGACAAGT	1	22	45.45	60.74
SARSCoV_1200_29_LEFT	TGAGGGAGCCTTGAATACACC A	1	22	50	61.1
SARSCoV_1200_29_RIGHT	TAGGCAGCTCTCCCTAGCATTG	1	22	54.55	61.61
SARSCoV_1200_2_LEFT	CCATAATCAAGACTATTCAACC AAGGGT	2	28	39.29	61.27
SARSCoV_1200_2_RIGHT	ACAGGTGACAATTTGTCCACCG	2	22	50	61.33
SARSCoV_1200_4_LEFT	GGAATTTGGTGCCACTTCTGCT	2	22	50	61.66
SARSCoV_1200_4_RIGHT	CCTGACCCGGGTAAGTGGTTAT	2	22	54.55	61.49
SARSCoV_1200_6_LEFT	ACTTCTATTAATGGGCAGATA ACAACCTG	2	29	34.48	60.18

SARSCoV_1200_6_RIGHT	GATTATCCATTCCCTGCGCGTC	2	22	54.55	61.75
SARSCoV_1200_8_LEFT	CAATCATGCAATTGTTTTTCAGC TATTTTG	2	30	30	60.39
SARSCoV_1200_8_RIGHT	TGACTTTTTGCTACCTGCGCAT	2	22	45.45	61.39
SARSCoV_1200_10_LEFT	TTTACCAGGAGTTTTCTGTGGT GT	2	24	41.67	60.32
SARSCoV_1200_10_RIGHT	TGGGCCTCATAGCACATTGGTA	2	22	50	61.5
SARSCoV_1200_12_LEFT	ATGGTGCTAGGAGAGTGTGGA C	2	22	54.55	61.48
SARSCoV_1200_12_RIGHT	GGATTTCCACAATGCTGATGC	2	22	50	60.48
SARSCoV_1200_14_LEFT	ACAGGCACTAGTACTGATGTCG T	2	23	47.83	61.12
SARSCoV_1200_14_RIGHT	GTGCAGCTACTGAAAAGCACG T	2	22	50	61.94
SARSCoV_1200_16_LEFT	ACAACACAGACTTTATGAGTGT CTCT	2	26	38.46	60.18
SARSCoV_1200_16_RIGHT	CTCTGTCAGACAGCACTTCACG	2	22	54.55	61.17
SARSCoV_1200_18_LEFT	GCACATAAAGACAAATCAGCTC AATGC	2	27	40.74	62.03
SARSCoV_1200_18_RIGHT	TGTCTGAAGCAGTGGAAAAGC A	2	22	45.45	60.68
SARSCoV_1200_20_LEFT	ACAATTTGATACTTATAACCTCT GGAACAC	2	30	33.33	60.15
SARSCoV_1200_20_RIGHT	GATTAGGCATAGCAACACCCG G	2	22	54.55	61.39
SARSCoV_1200_22_LEFT	GTGATGTTCTTGTTAACAATA AACGAACA	2	30	33.33	61.44
SARSCoV_1200_22_RIGHT	AACAGATGCAAATCTGGTGCC G	2	22	50	62.03
SARSCoV_1200_24_LEFT	GCTGAACATGTCAACAACATCAT ATGA	2	26	38.46	60.13
SARSCoV_1200_24_RIGHT	ATGAGGTGCTGACTGAGGGAA G	2	22	54.55	61.74
SARSCoV_1200_26_LEFT	GCCTTGAAGCCCCTTTTCTCTA	2	22	50	60.29

SARSCoV_1200_26_RIGHT	AATGACCACATGGAACGCGTA C	2	22	50	61.5
SARSCoV_1200_28_LEFT	TTTGTGCTTTTTAGCCTTTCTGC T	2	24	37.5	60.14
SARSCoV_1200_28_RIGHT	GTTTGGCCTTGTTGTTGTTGGC	2	22	50	61.82